

anova p-value: 9.789e-05

fdr-adjusted p-value: 0.016837

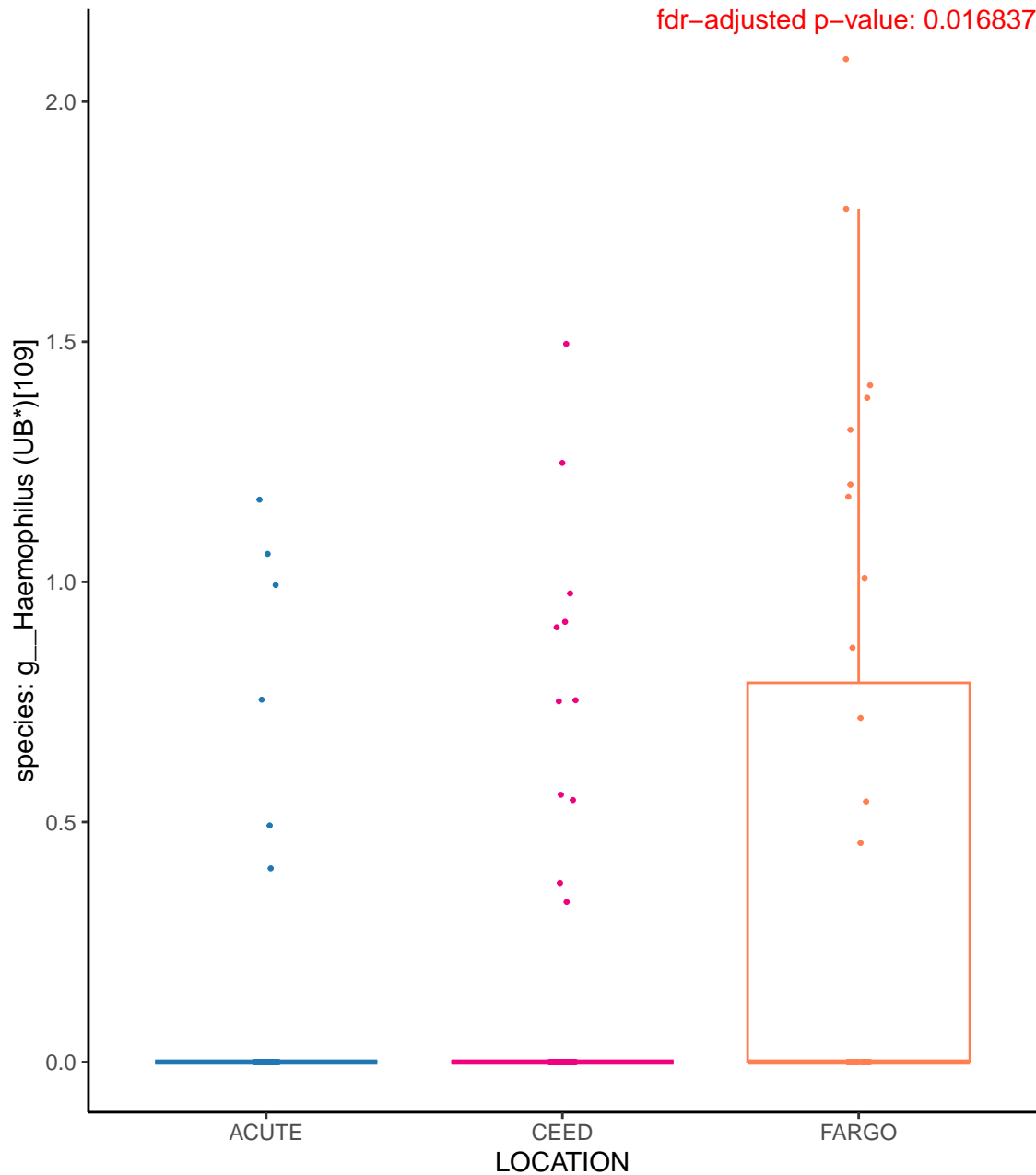
species: g__Haemophilus (UB*)[109]

LOCATION

ACUTE

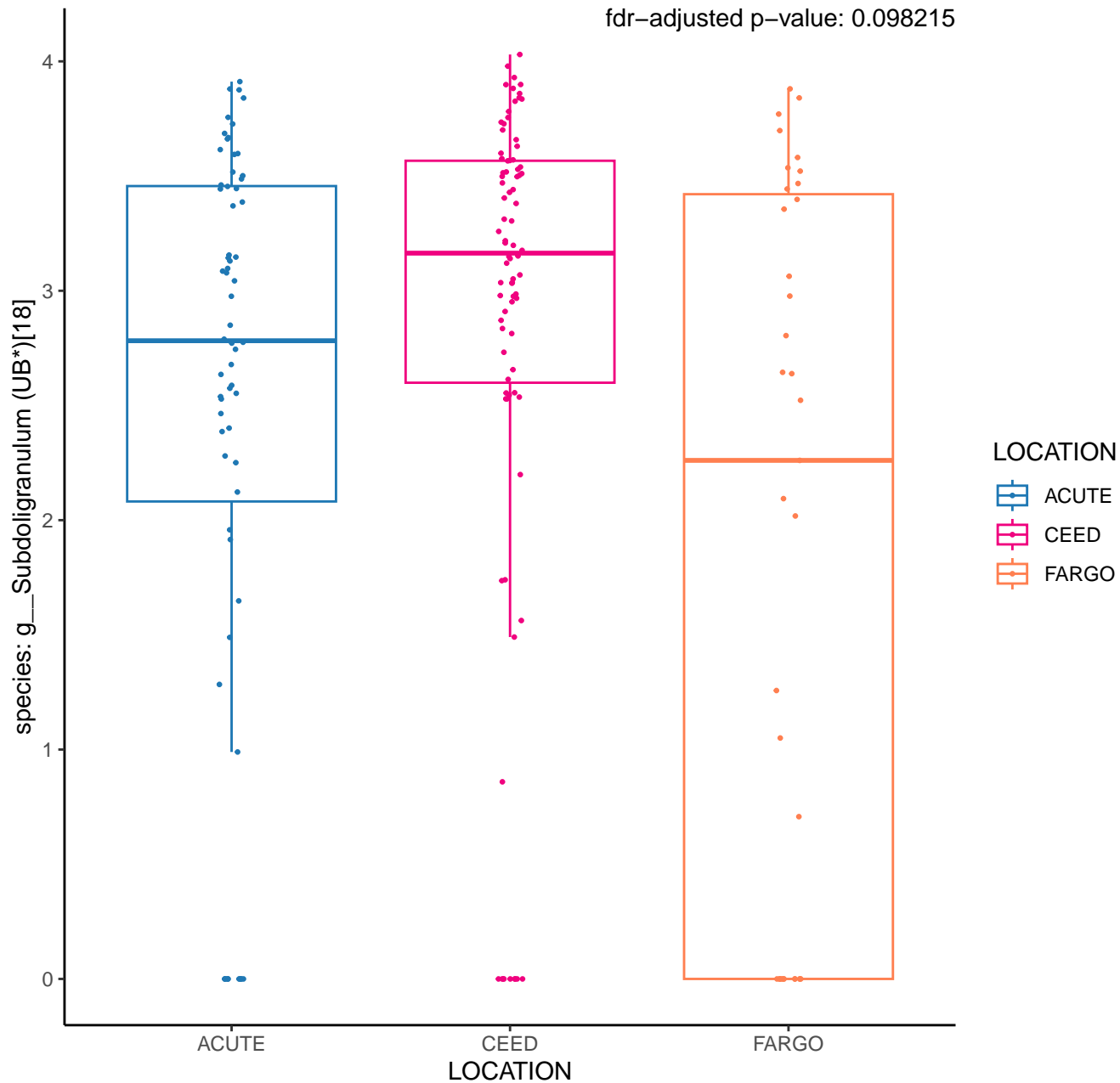
CEED

FARGO



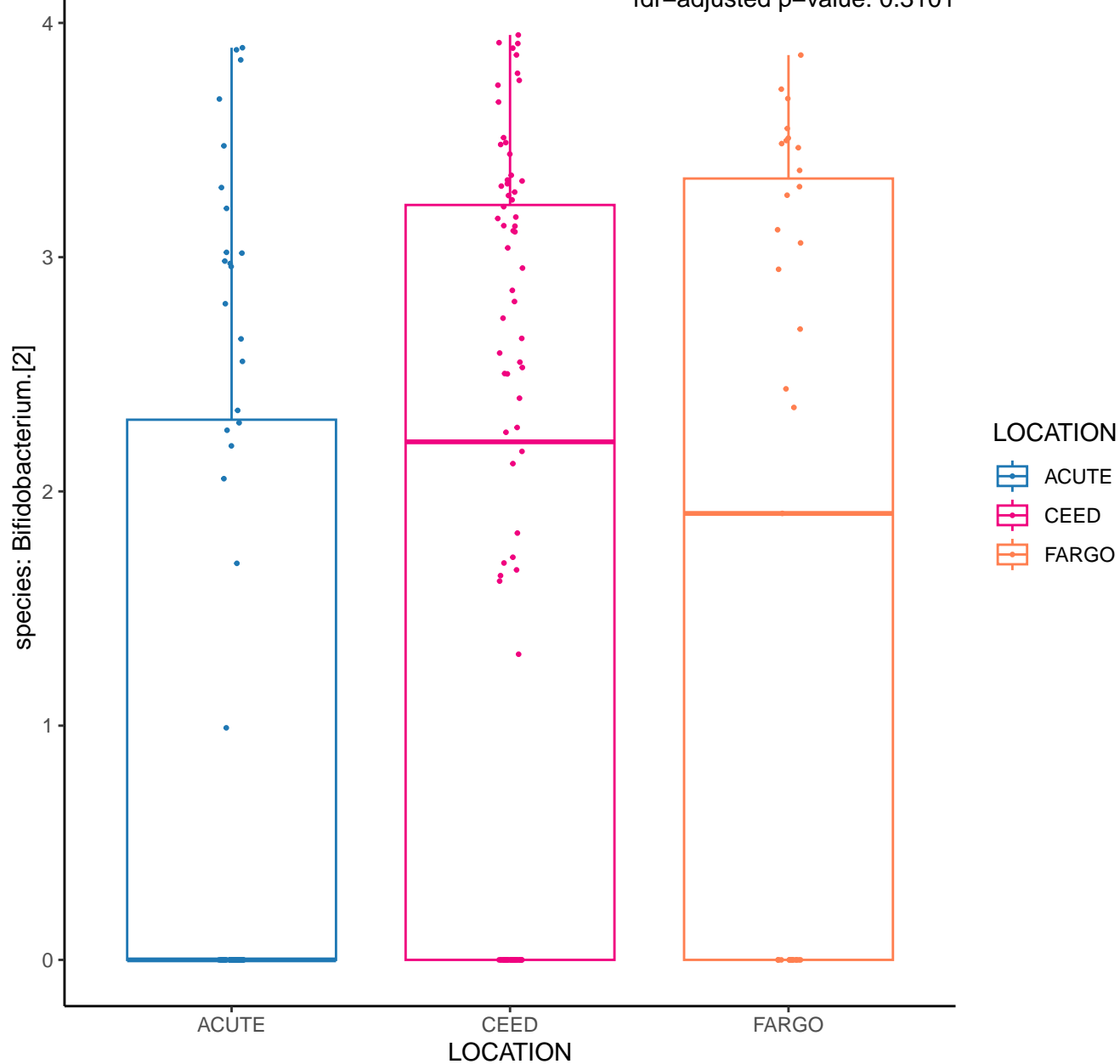
anova p-value: 0.001142

fdr-adjusted p-value: 0.098215



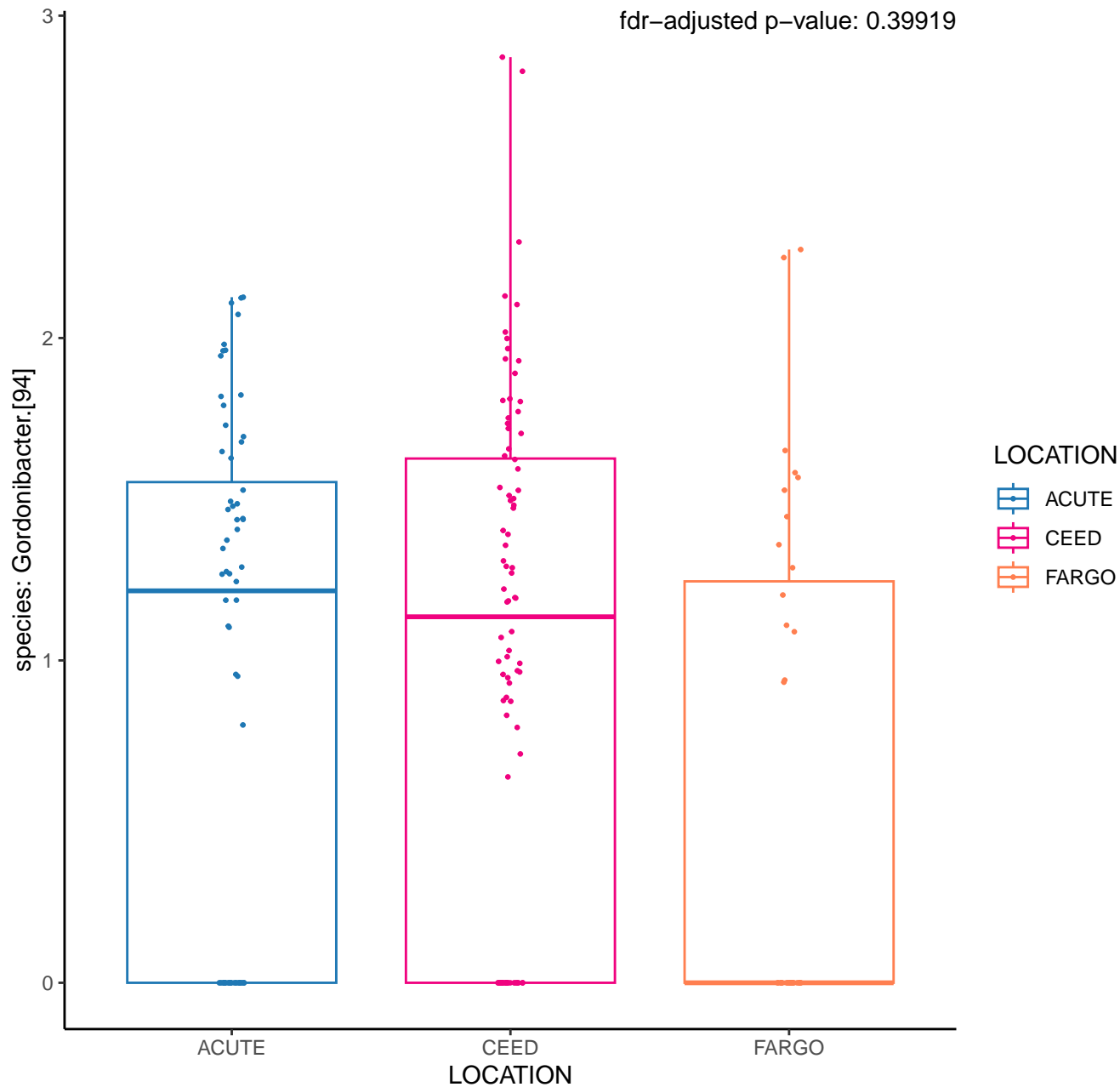
anova p-value: 0.0054086

fdr-adjusted p-value: 0.3101



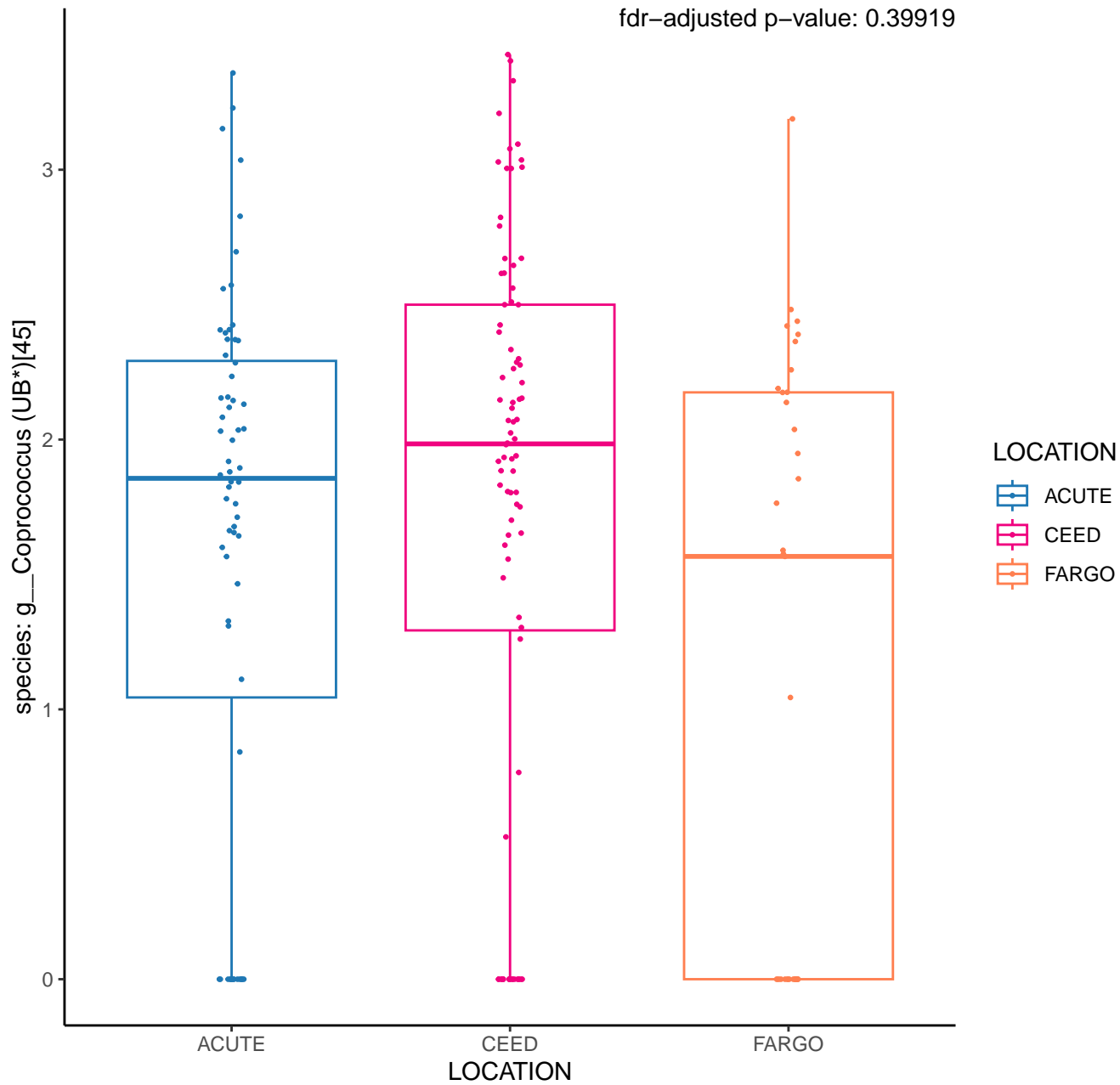
anova p-value: 0.015424

fdr-adjusted p-value: 0.39919



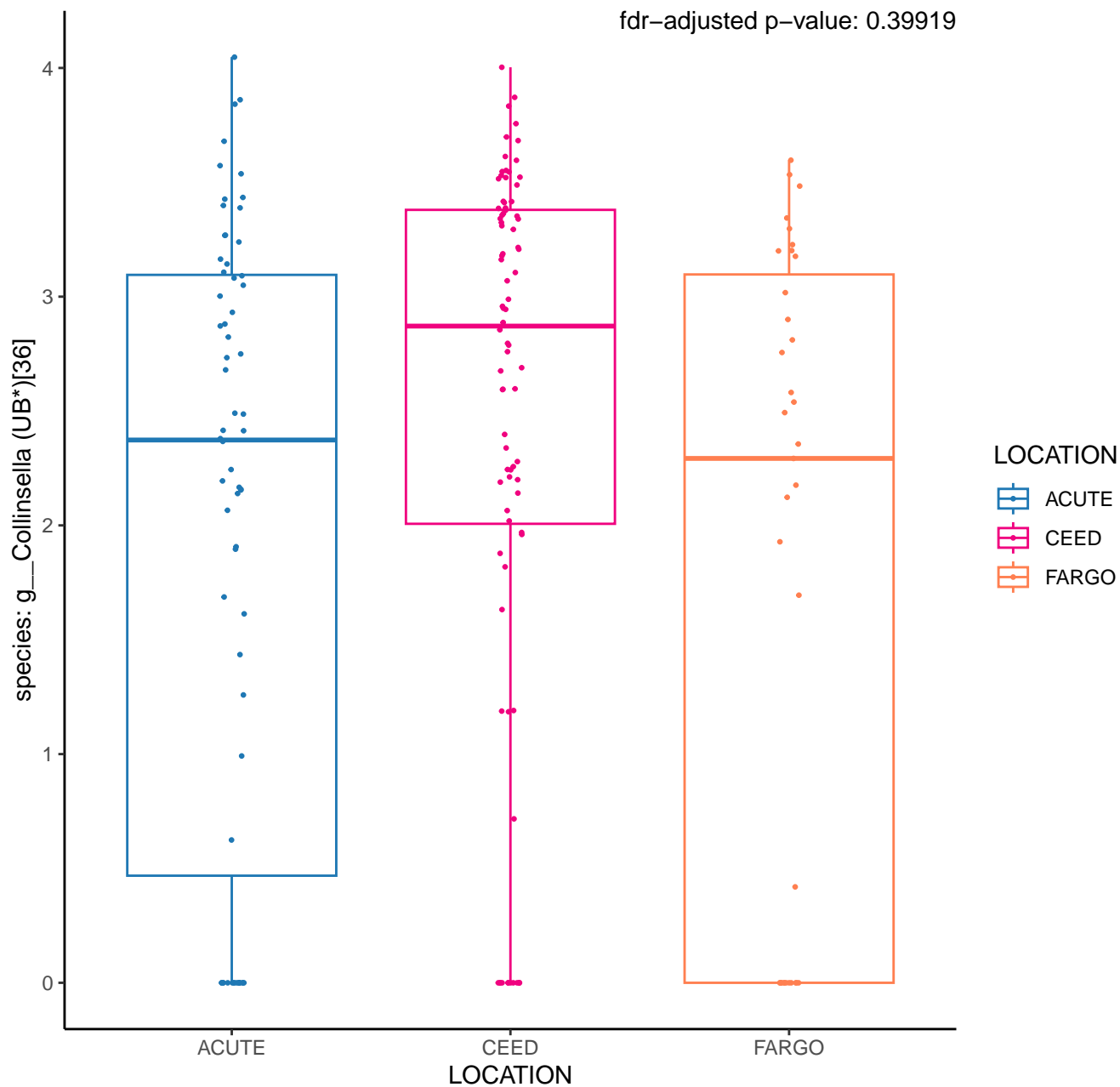
anova p-value: 0.016398

fdr-adjusted p-value: 0.39919



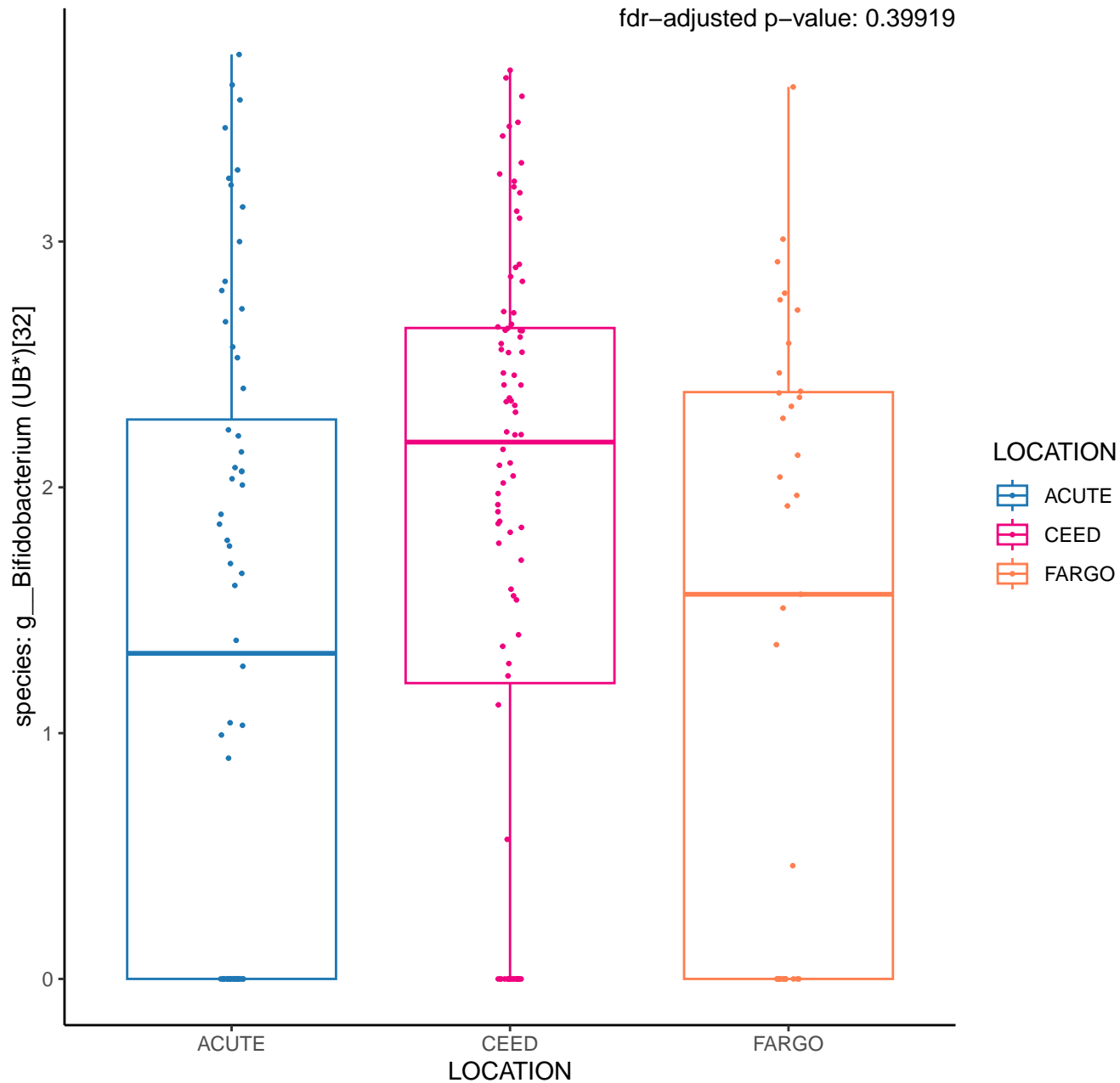
anova p-value: 0.016563

fdr-adjusted p-value: 0.39919



anova p-value: 0.017019

fdr-adjusted p-value: 0.39919



anova p-value: 0.018567

fdr-adjusted p-value: 0.39919

species: *Erysipelatoclostridium*. [77]

LOCATION

ACUTE

CEED

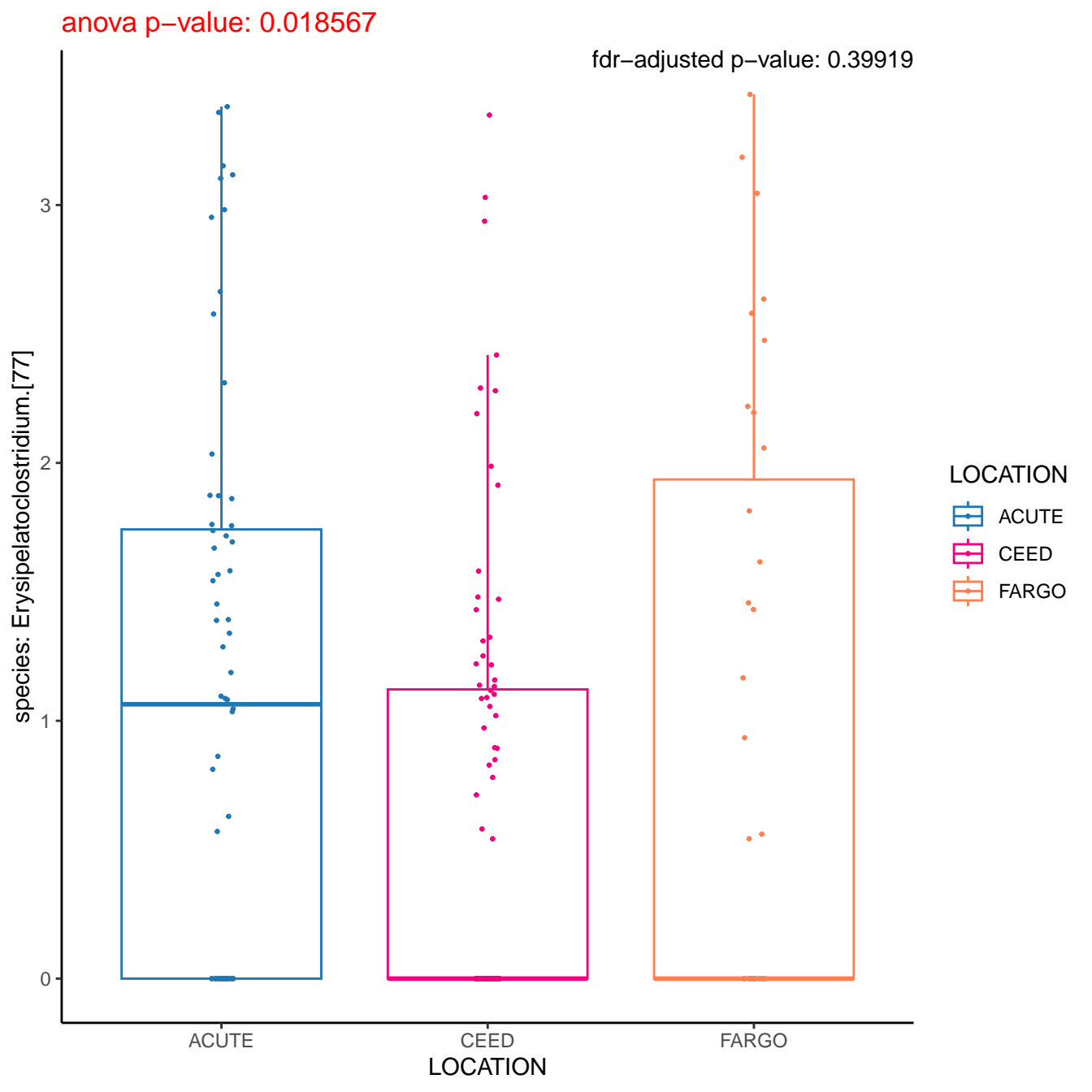
FARGO

ACUTE

CEED

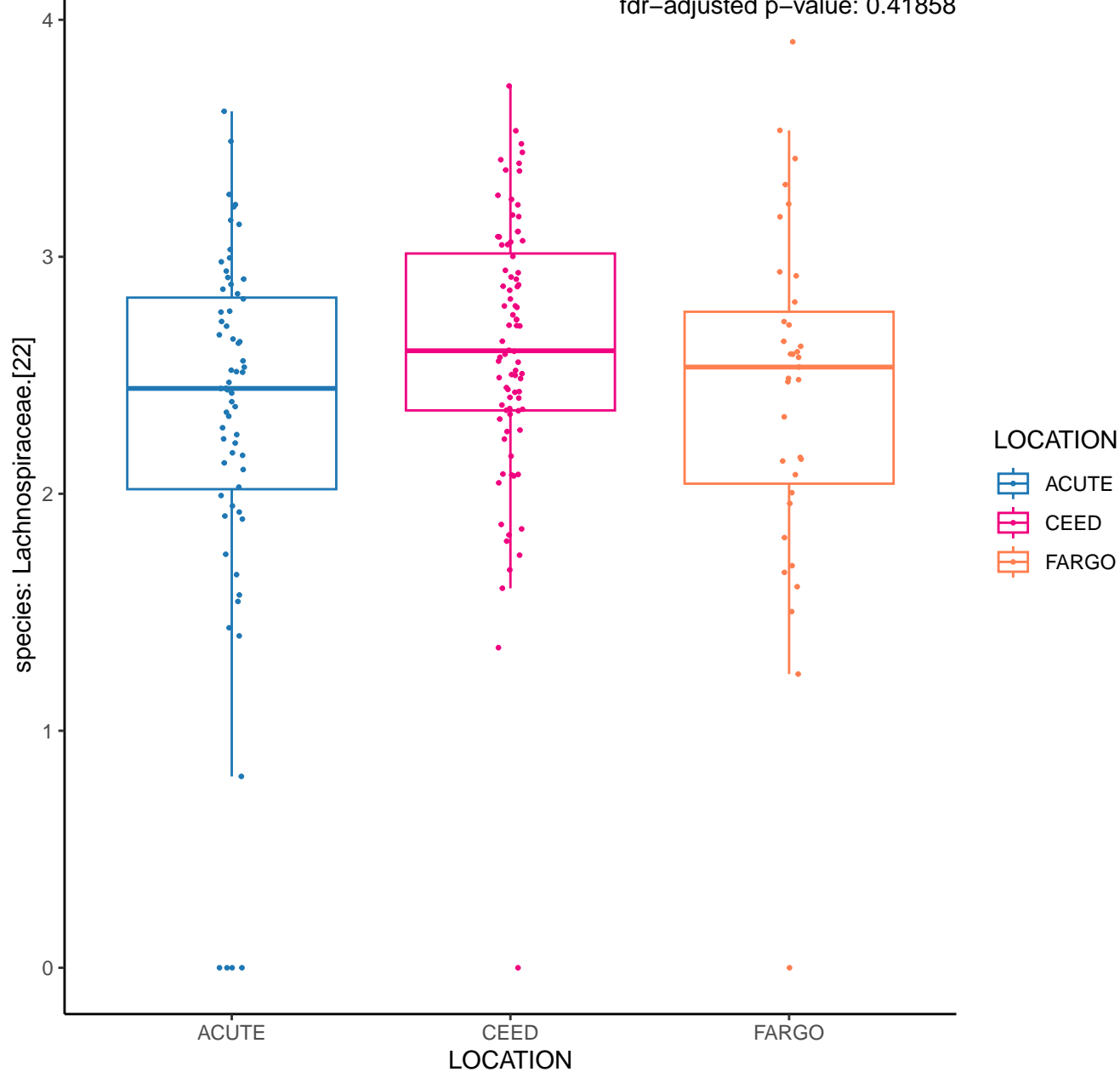
FARGO

LOCATION



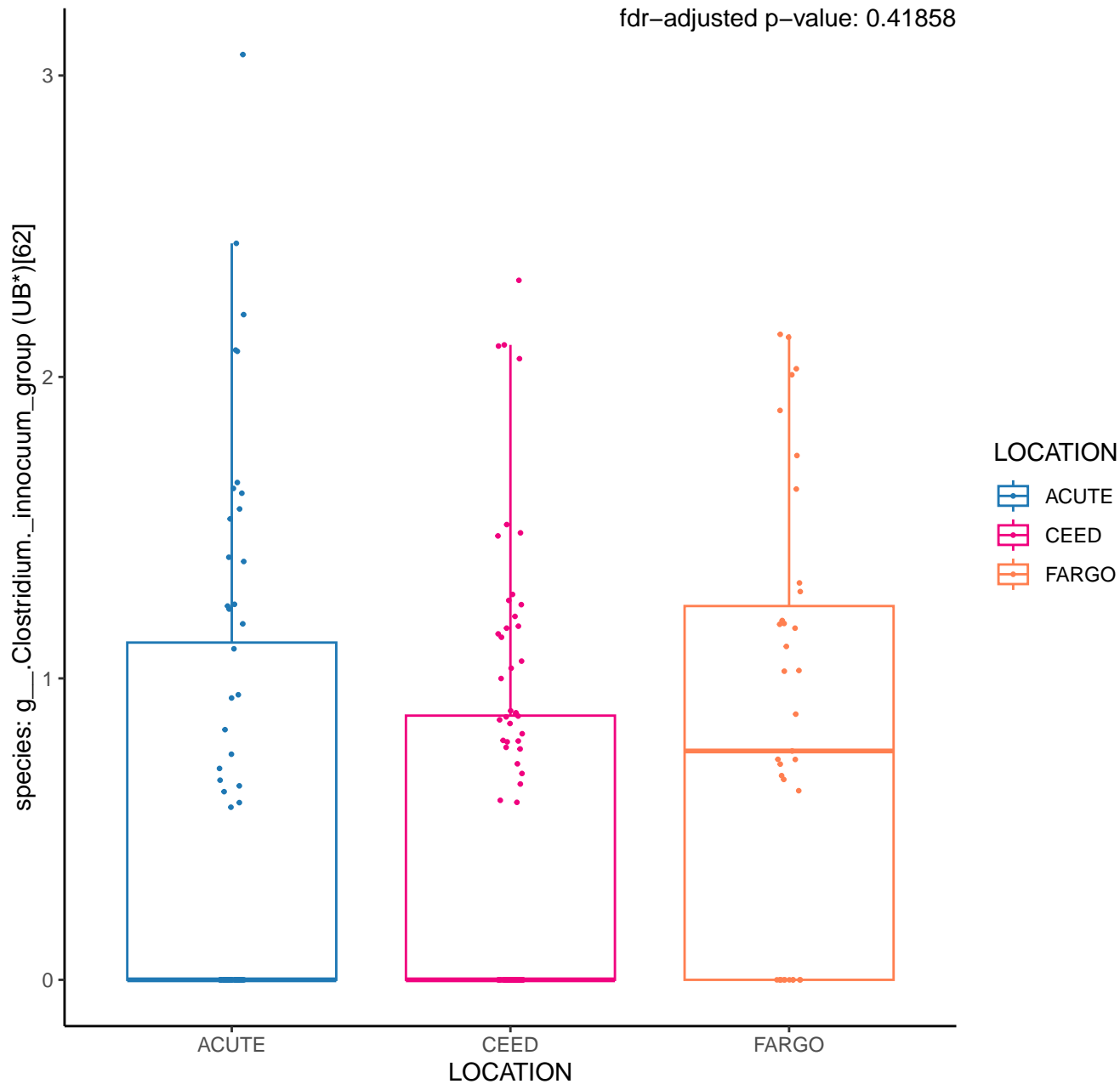
anova p-value: 0.027325

fdr-adjusted p-value: 0.41858



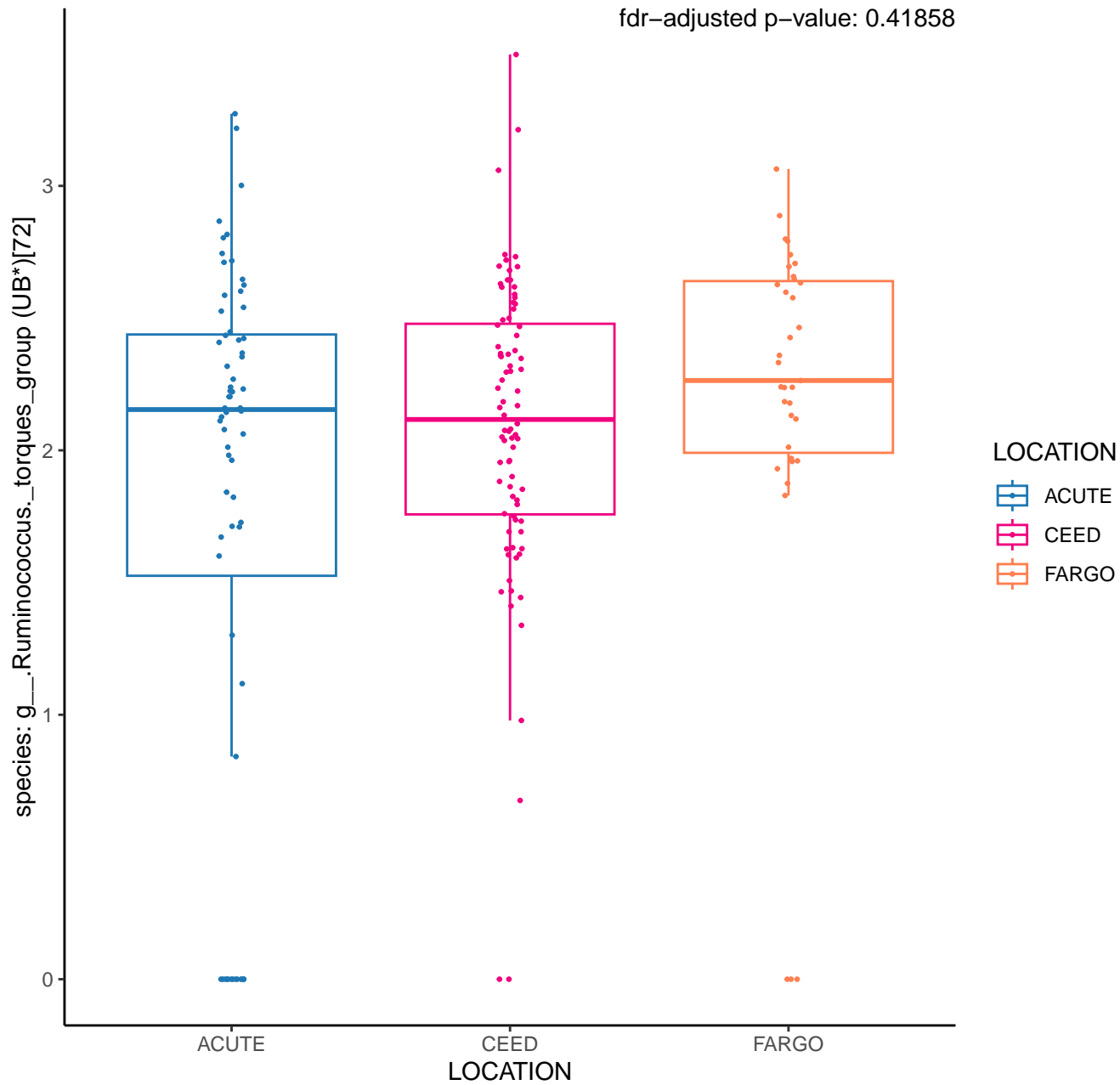
anova p-value: 0.028161

fdr-adjusted p-value: 0.41858



anova p-value: 0.028958

fdr-adjusted p-value: 0.41858

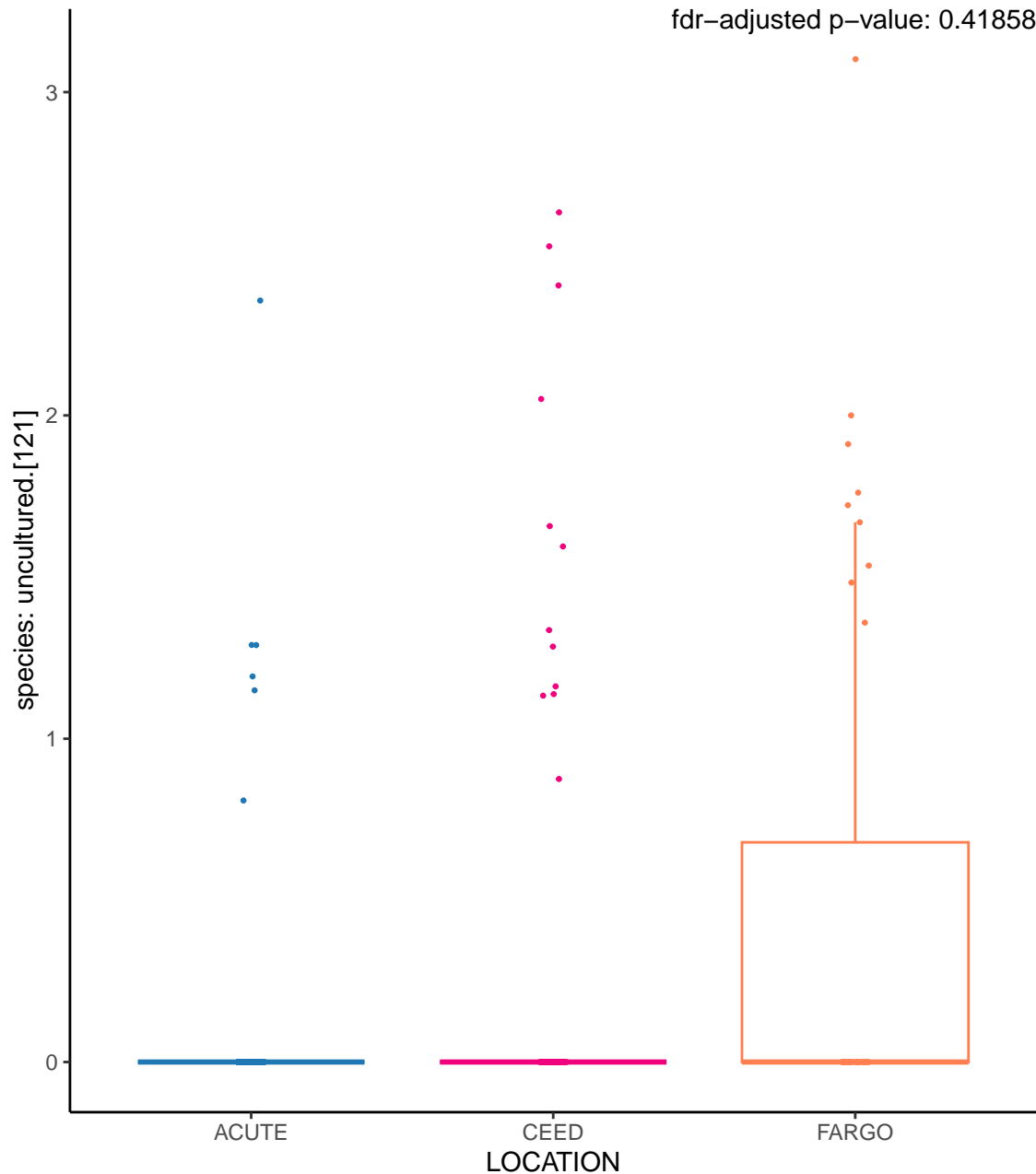


anova p-value: 0.029203

fdr-adjusted p-value: 0.41858

species: uncultured.[121]

LOCATION



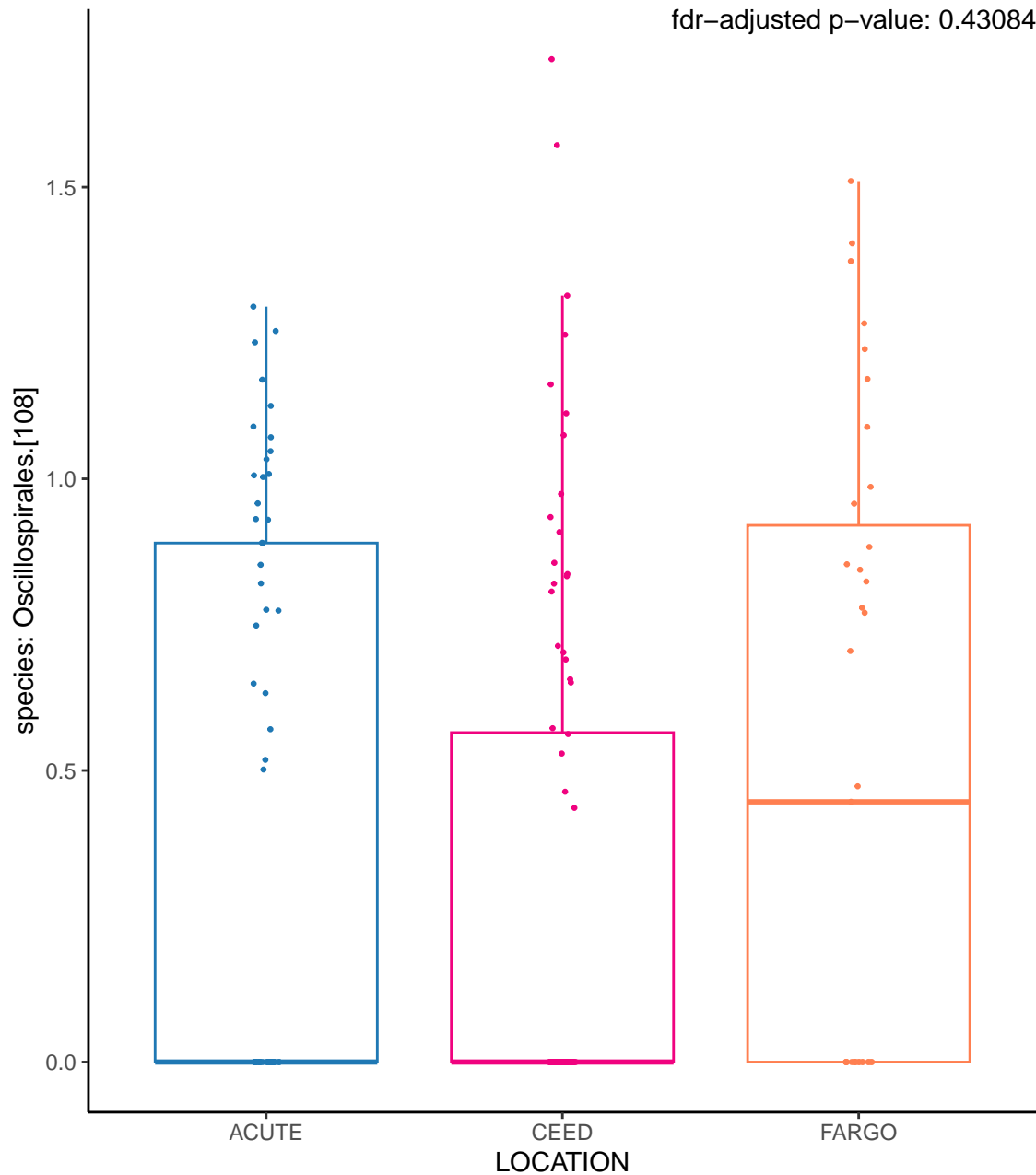
anova p-value: 0.036943

fdr-adjusted p-value: 0.43084

species: Oscillospirales.[108]

LOCATION

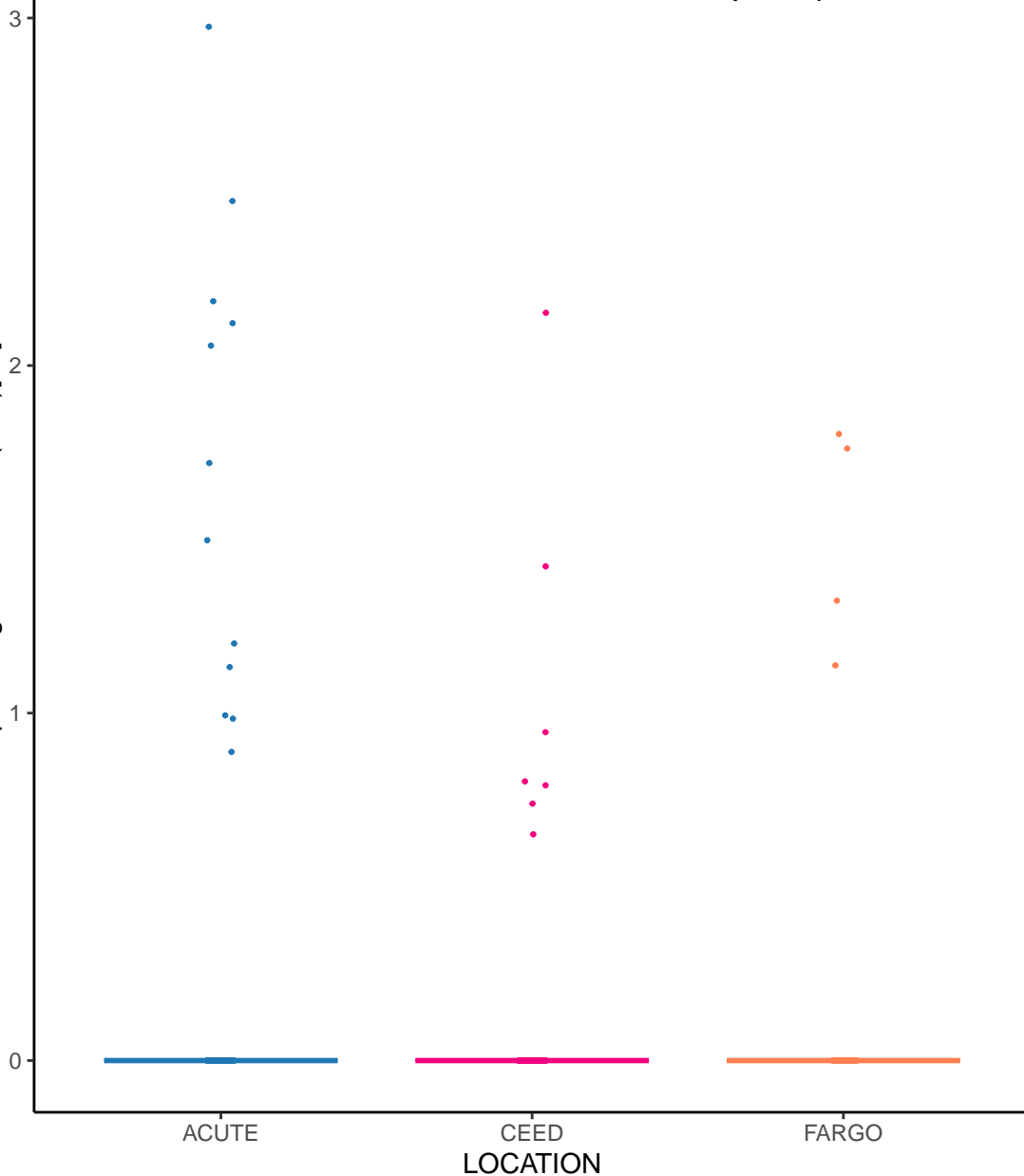
ACUTE
CEED
FARGO



anova p-value: 0.037035

fdr-adjusted p-value: 0.43084

species: g__Veillonella (UB*)[30]



LOCATION

ACUTE

CEED

FARGO

anova p-value: 0.041304

fdr-adjusted p-value: 0.43084

species: g__Dialister (UB*)[12]

LOCATION

ACUTE

CEED

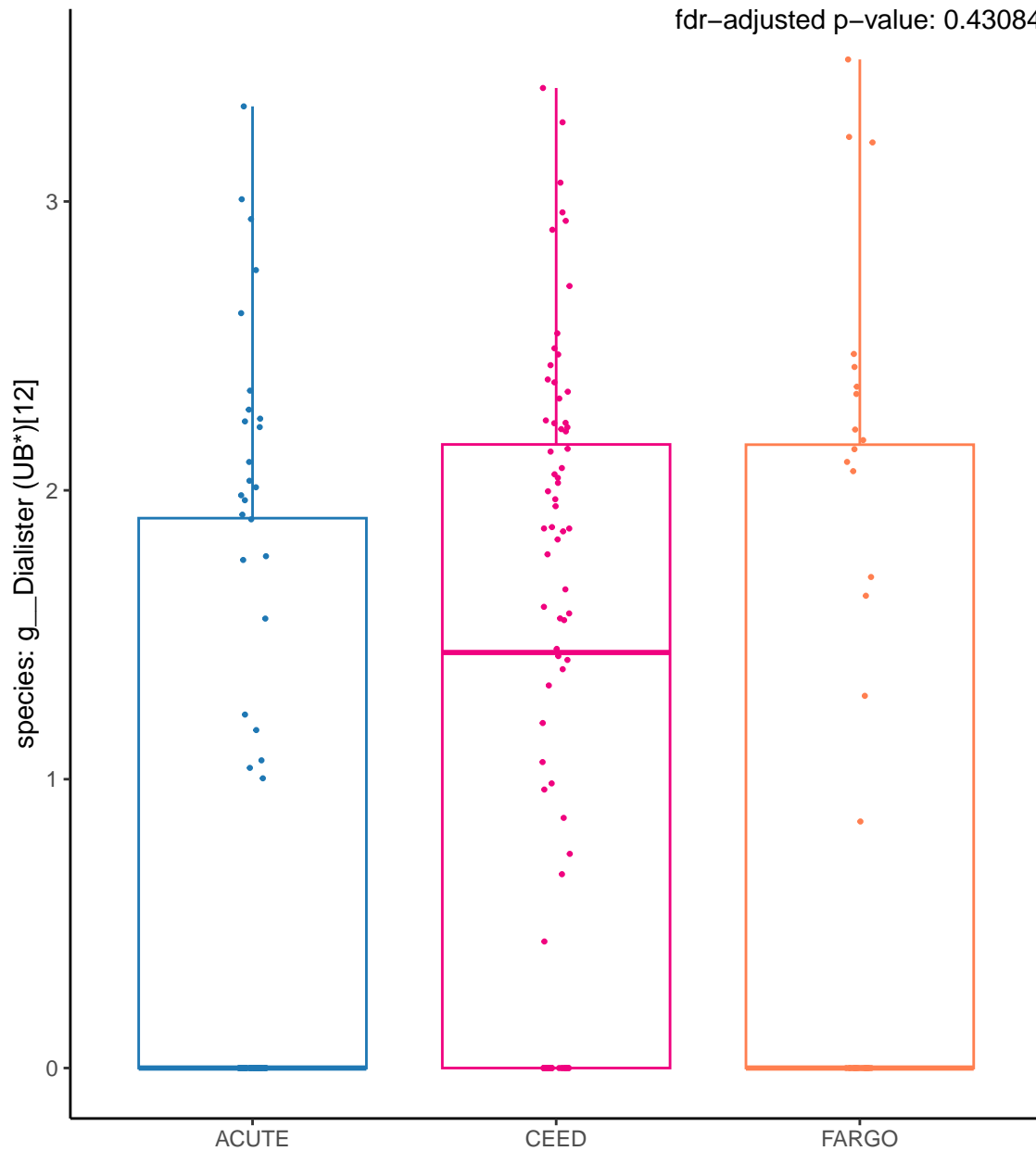
FARGO

ACUTE

CEED

FARGO

LOCATION



anova p-value: 0.042279

fdr-adjusted p-value: 0.43084

species: g__Romboutsia (UB*)[90]

LOCATION

ACUTE

CEED

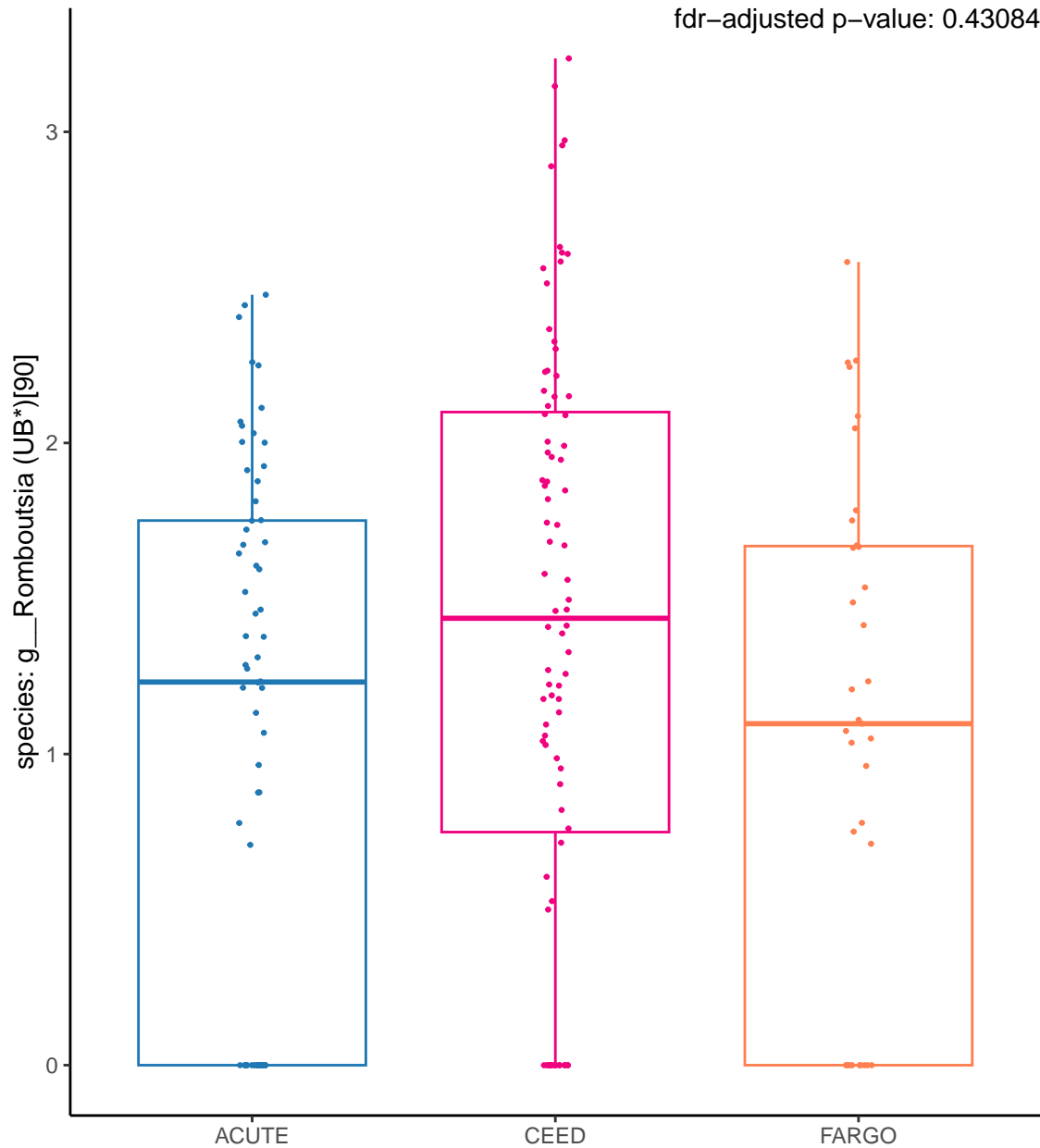
FARGO

ACUTE

CEED

FARGO

LOCATION



anova p-value: 0.045093

fdr-adjusted p-value: 0.43084

species: Streptococcus.[89]

LOCATION

ACUTE

CEED

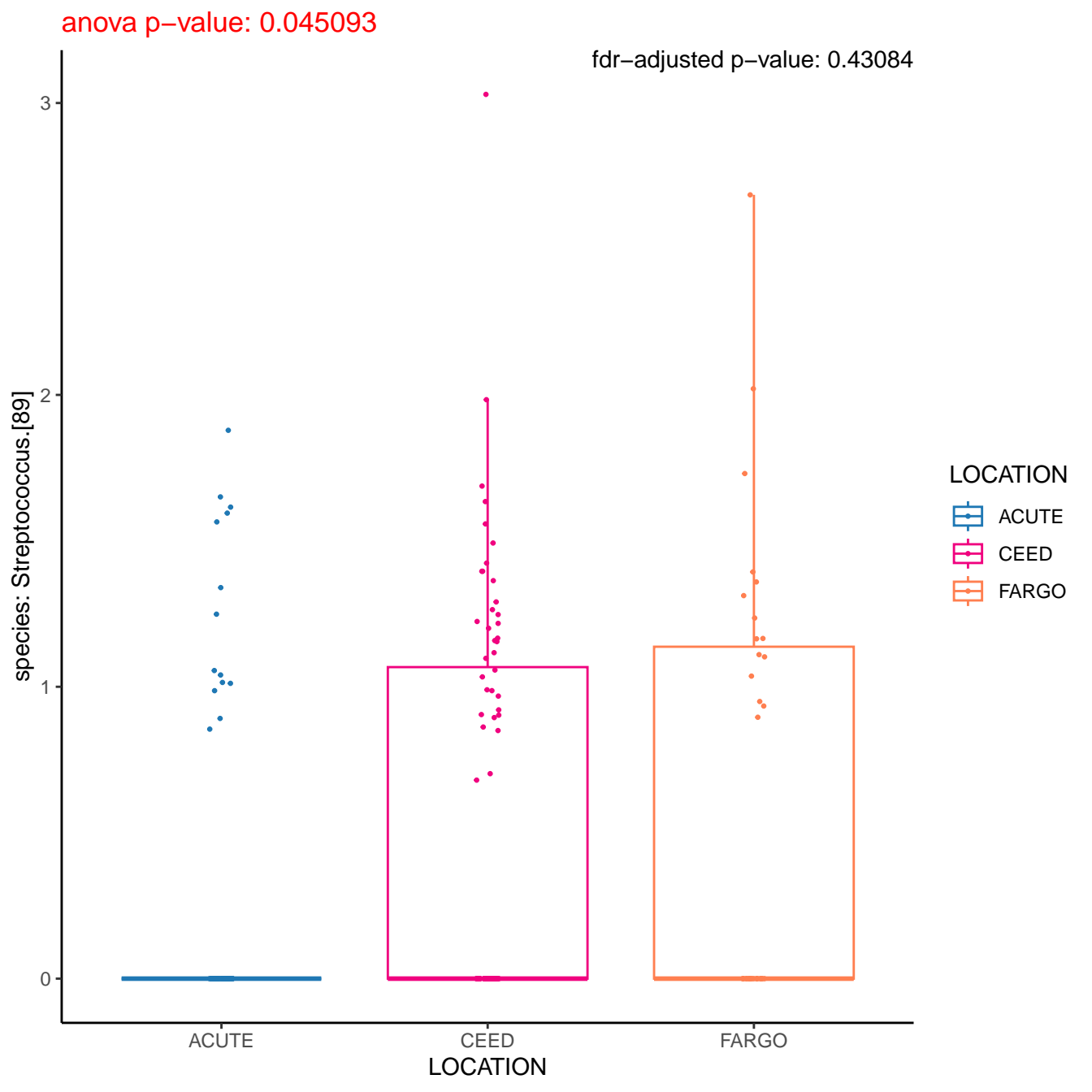
FARGO

ACUTE

CEED

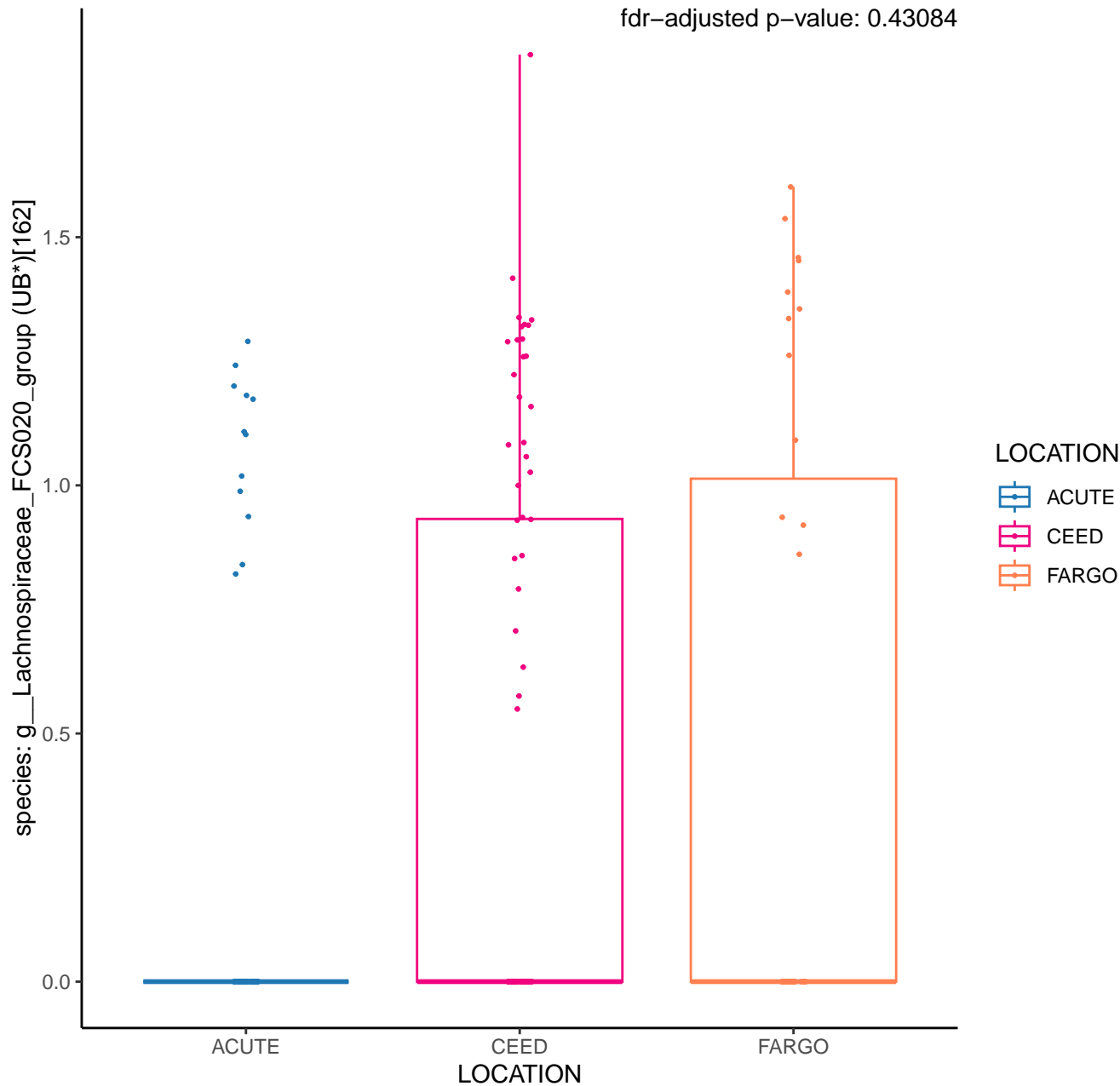
FARGO

LOCATION



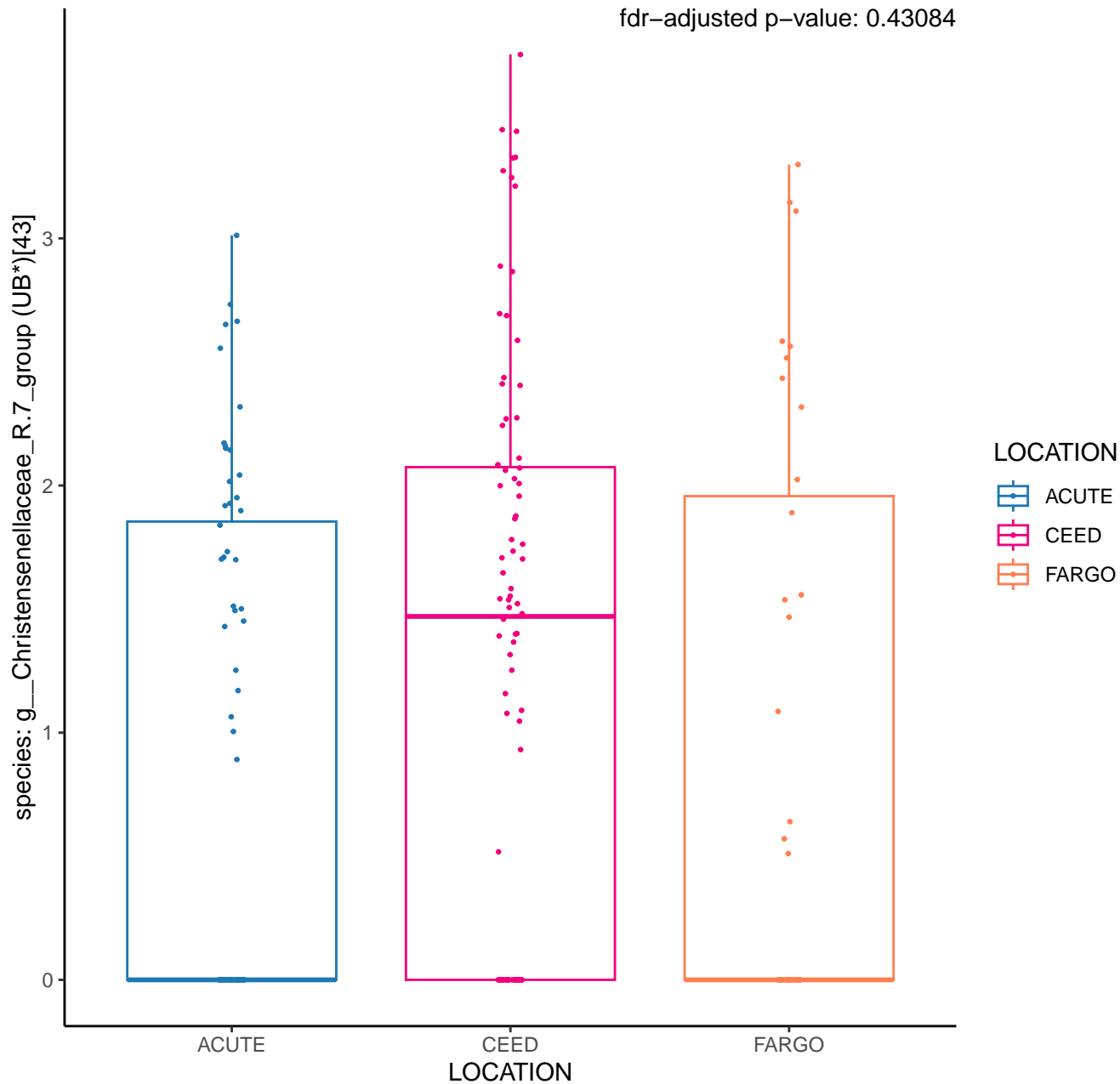
anova p-value: 0.04636

fdr-adjusted p-value: 0.43084



anova p-value: 0.047593

fdr-adjusted p-value: 0.43084

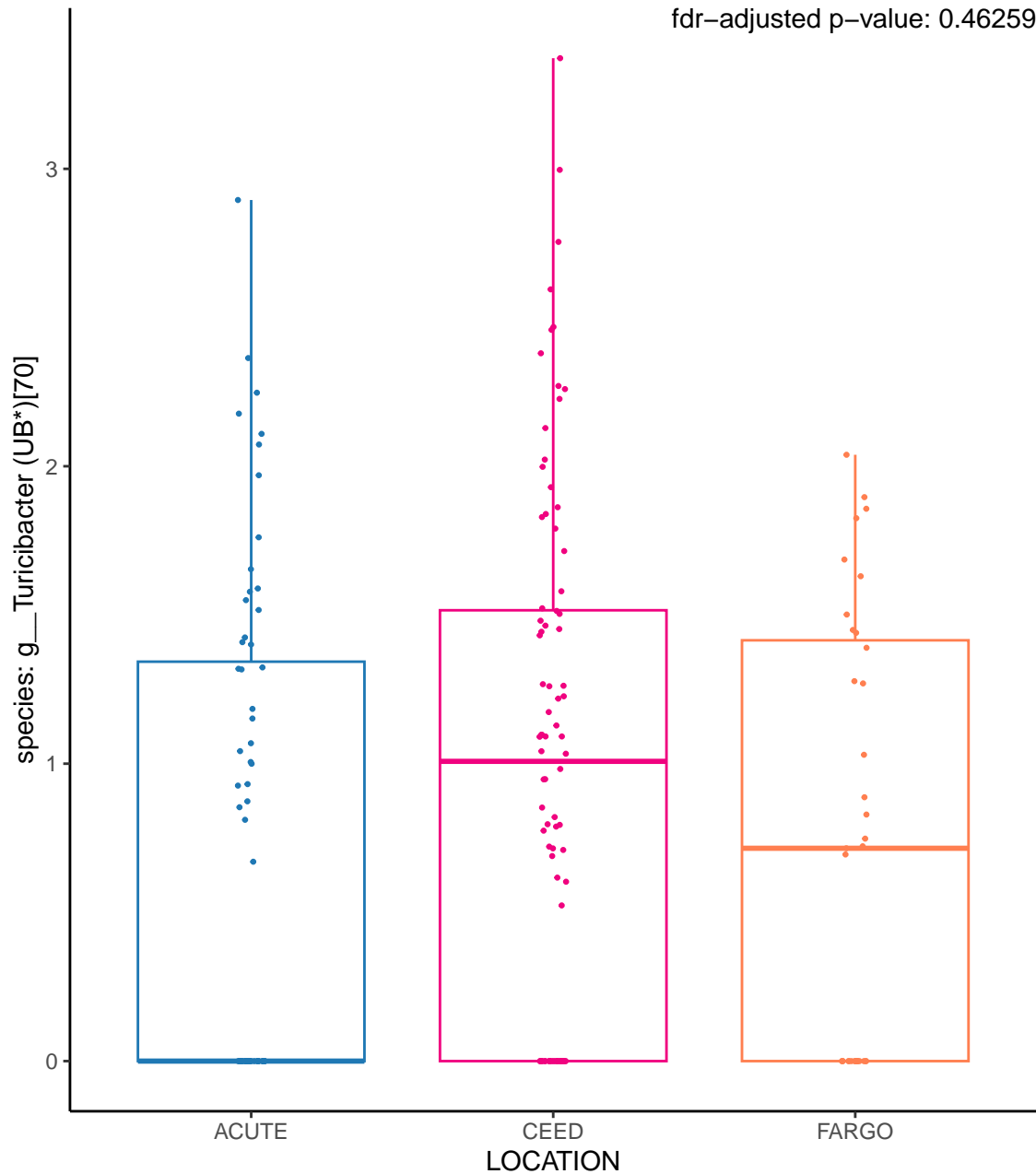


anova p-value: 0.054455

fdr-adjusted p-value: 0.46259

species: g__Turicibacter (UB*)[70]

LOCATION



anova p-value: 0.061076

fdr-adjusted p-value: 0.46259

species: *Clostridium_sensu_stricto_2*. [66]

LOCATION

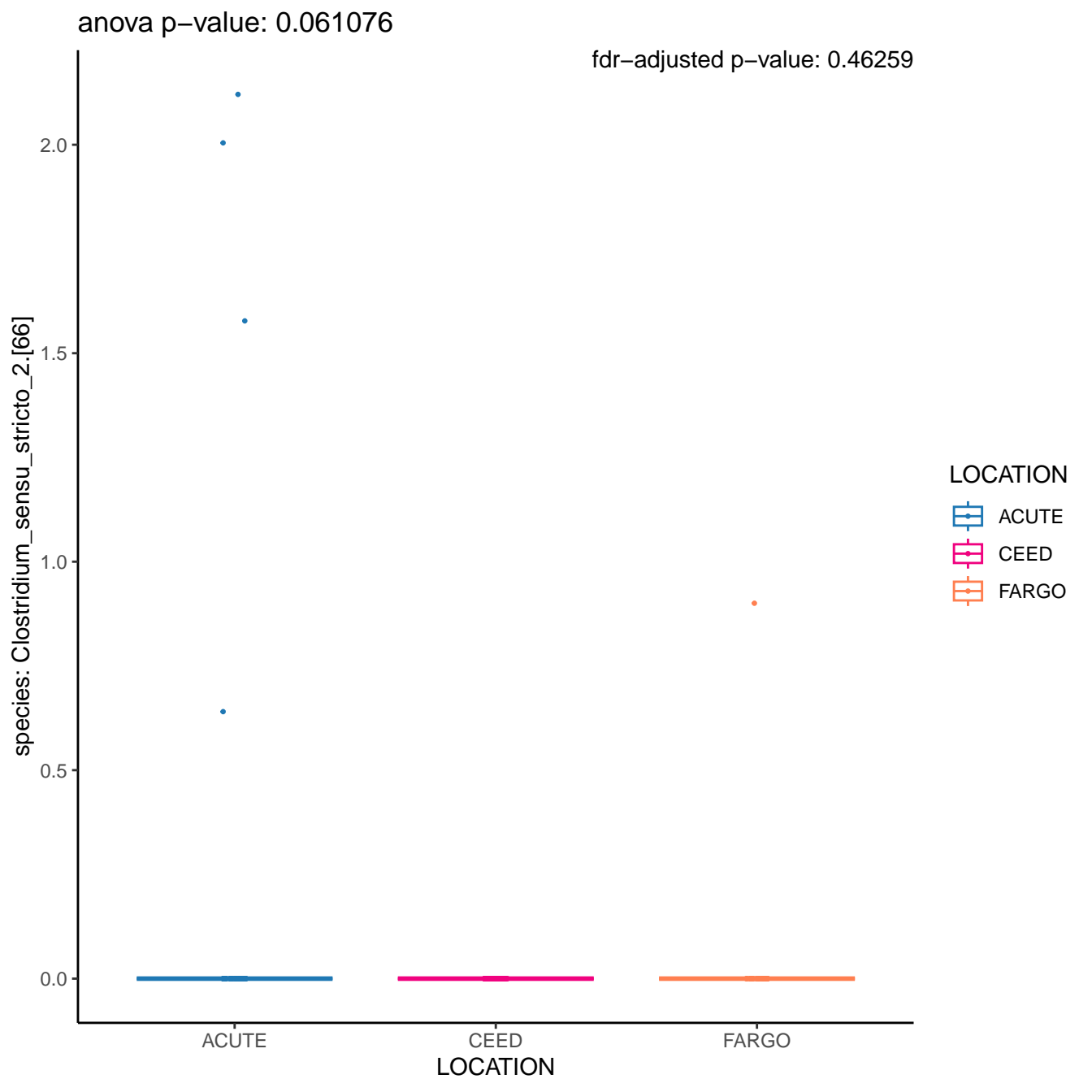
ACUTE
CEED
FARGO

ACUTE

CEED

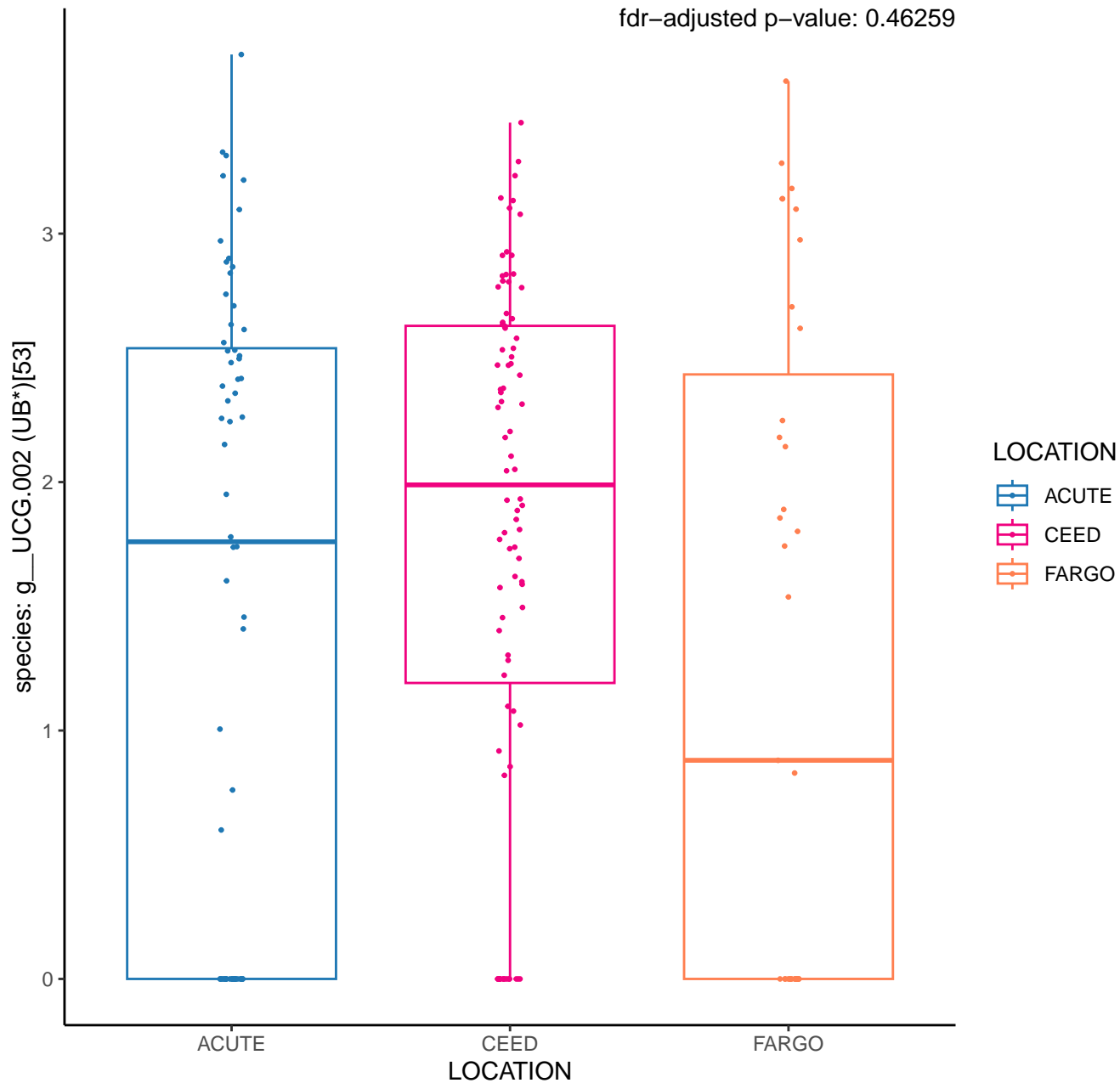
FARGO

LOCATION

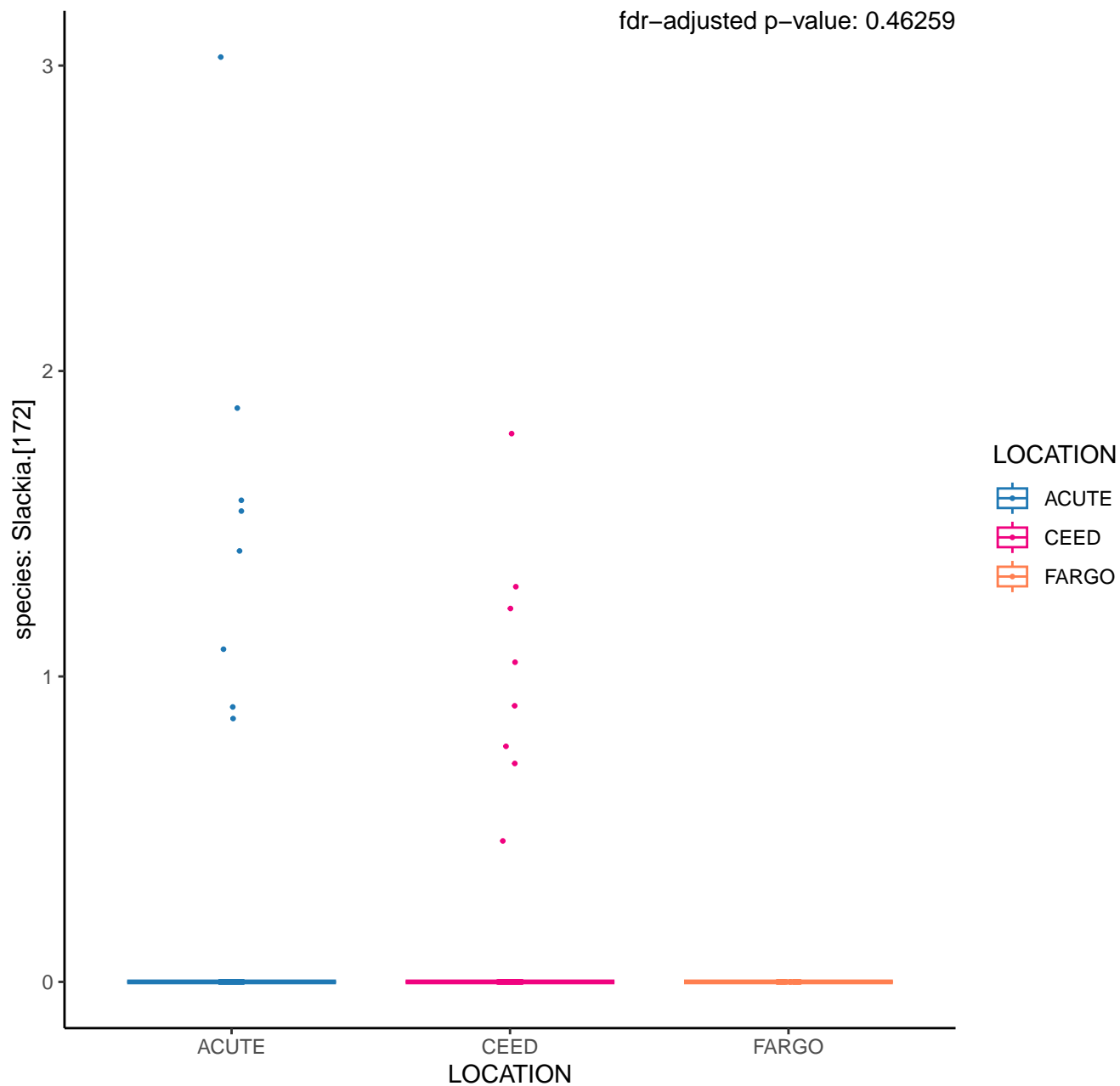


anova p-value: 0.061979

fdr-adjusted p-value: 0.46259

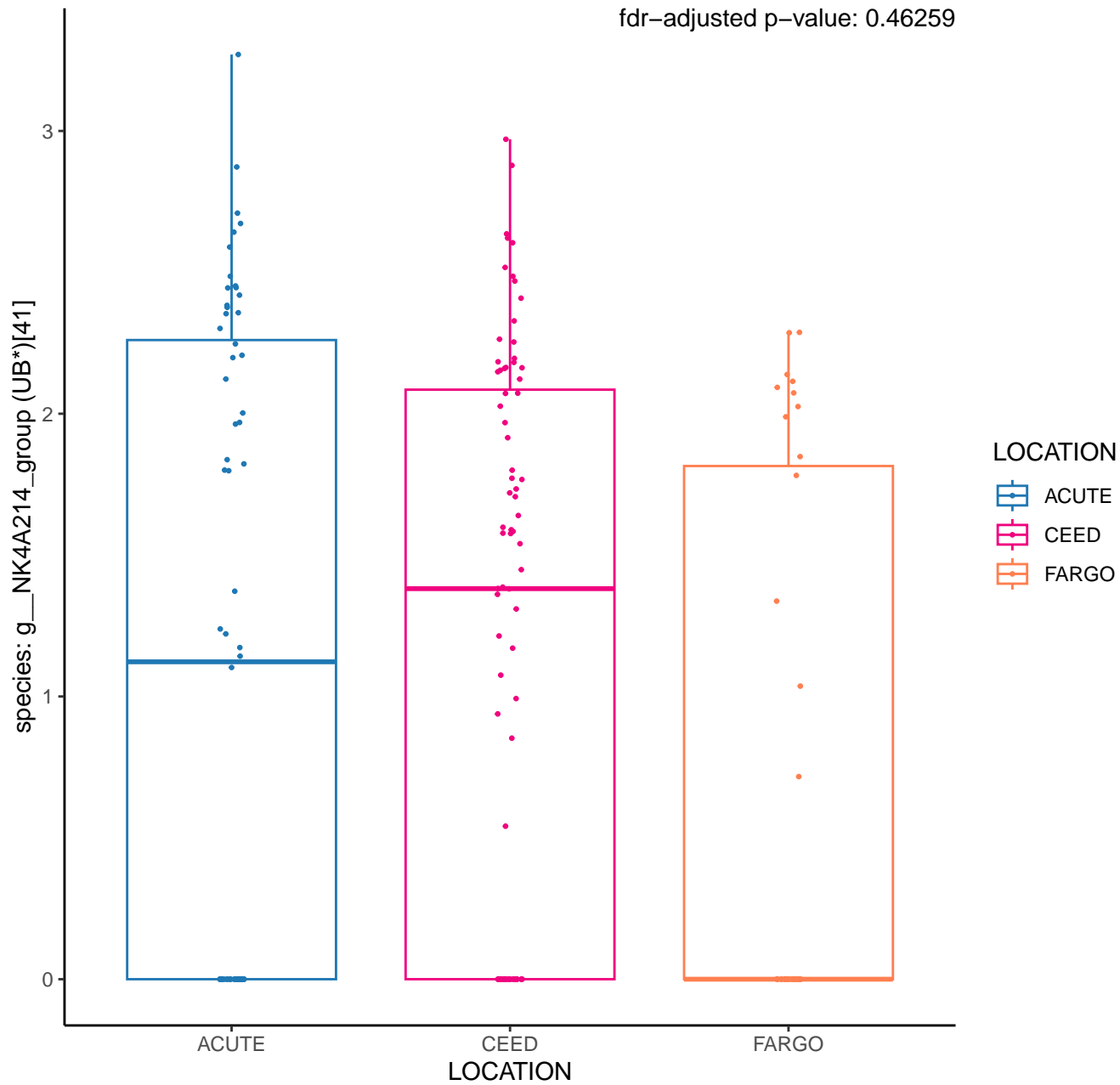


fdr-adjusted p-value: 0.46259



anova p-value: 0.071418

fdr-adjusted p-value: 0.46259



anova p-value: 0.076591

fdr-adjusted p-value: 0.46259

species: Anaerostipes.[92]

LOCATION

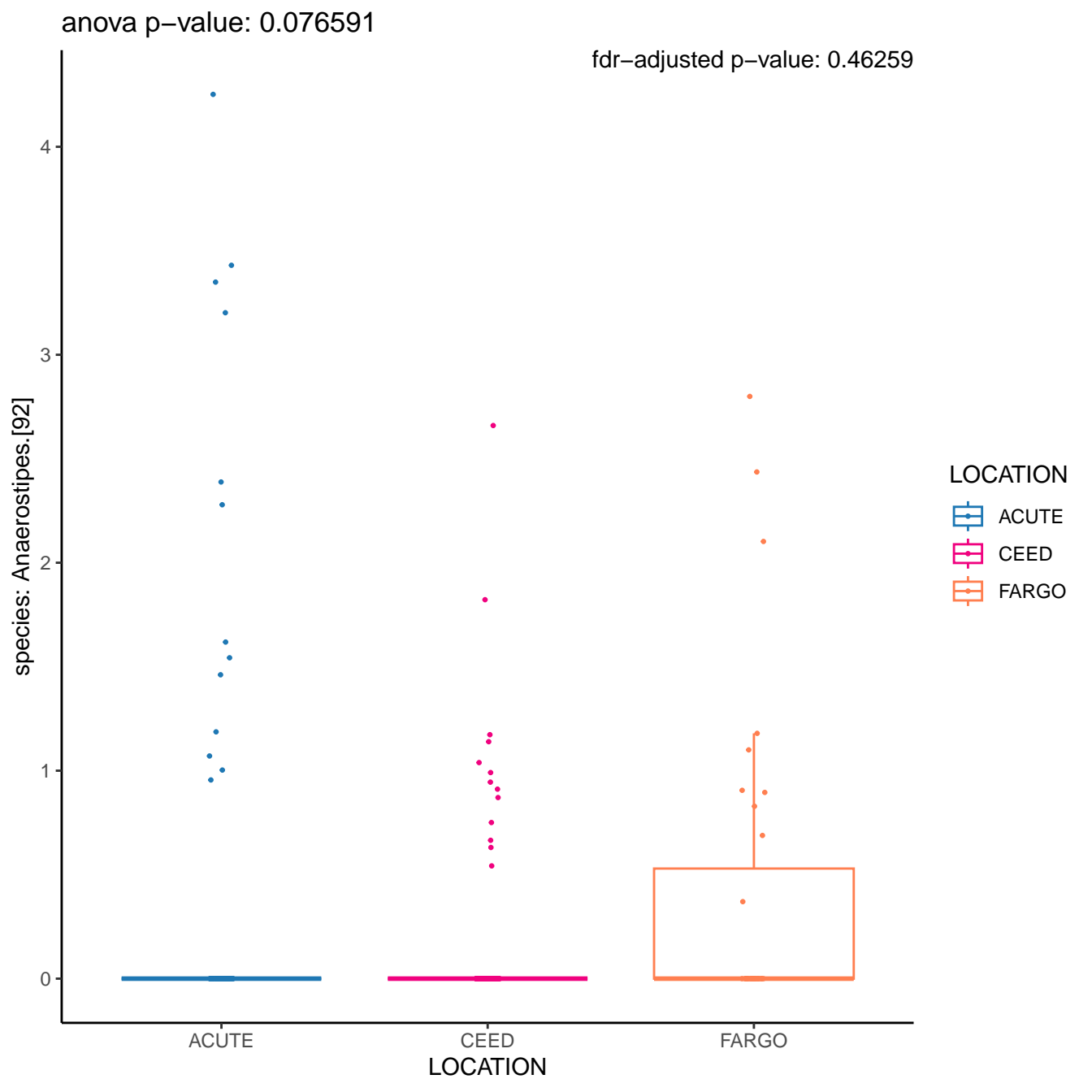


ACUTE

CEED

FARGO

LOCATION



anova p-value: 0.077288

fdr-adjusted p-value: 0.46259

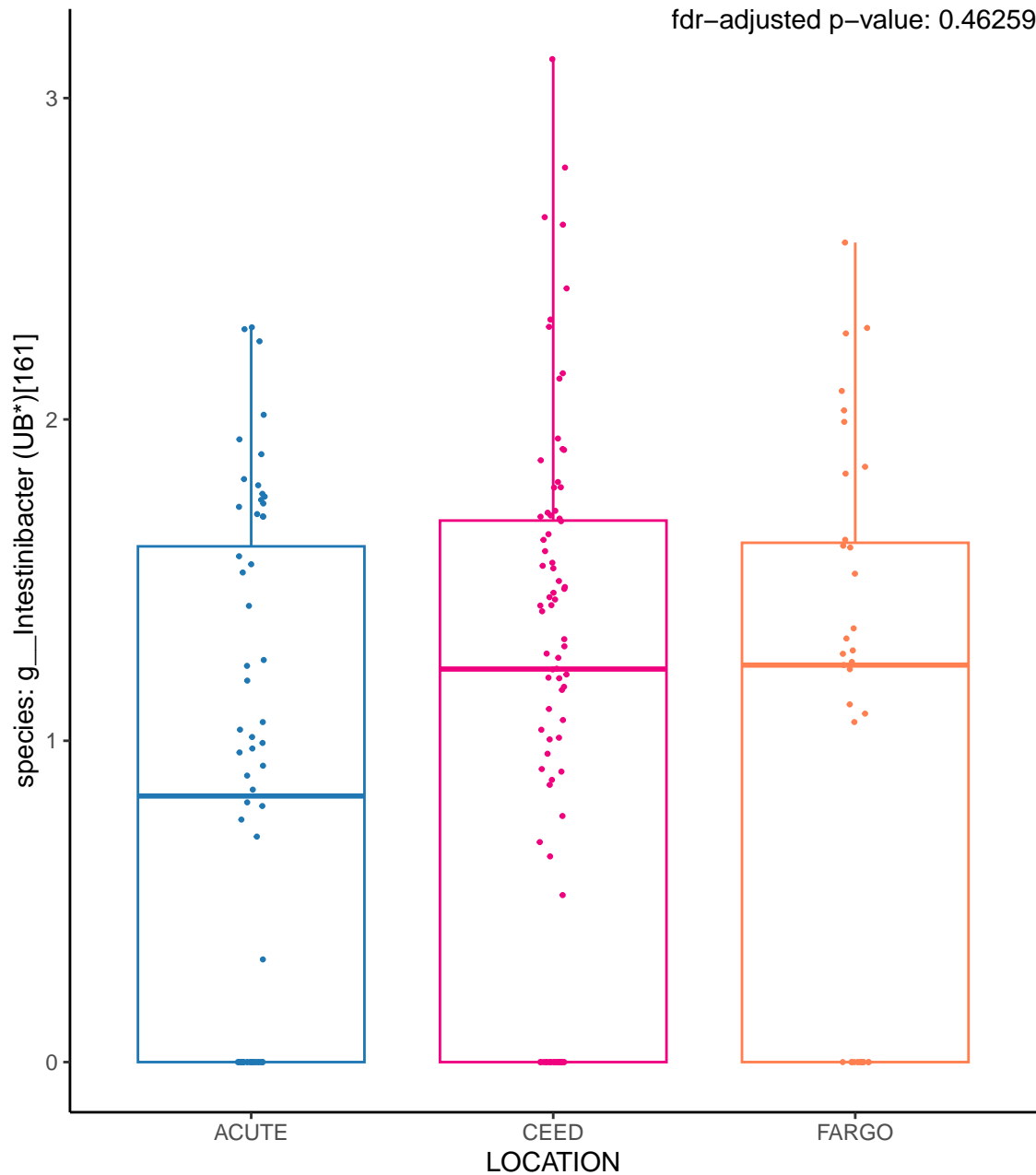
species: g__Intestinibacter (UB*)[161]

LOCATION

ACUTE

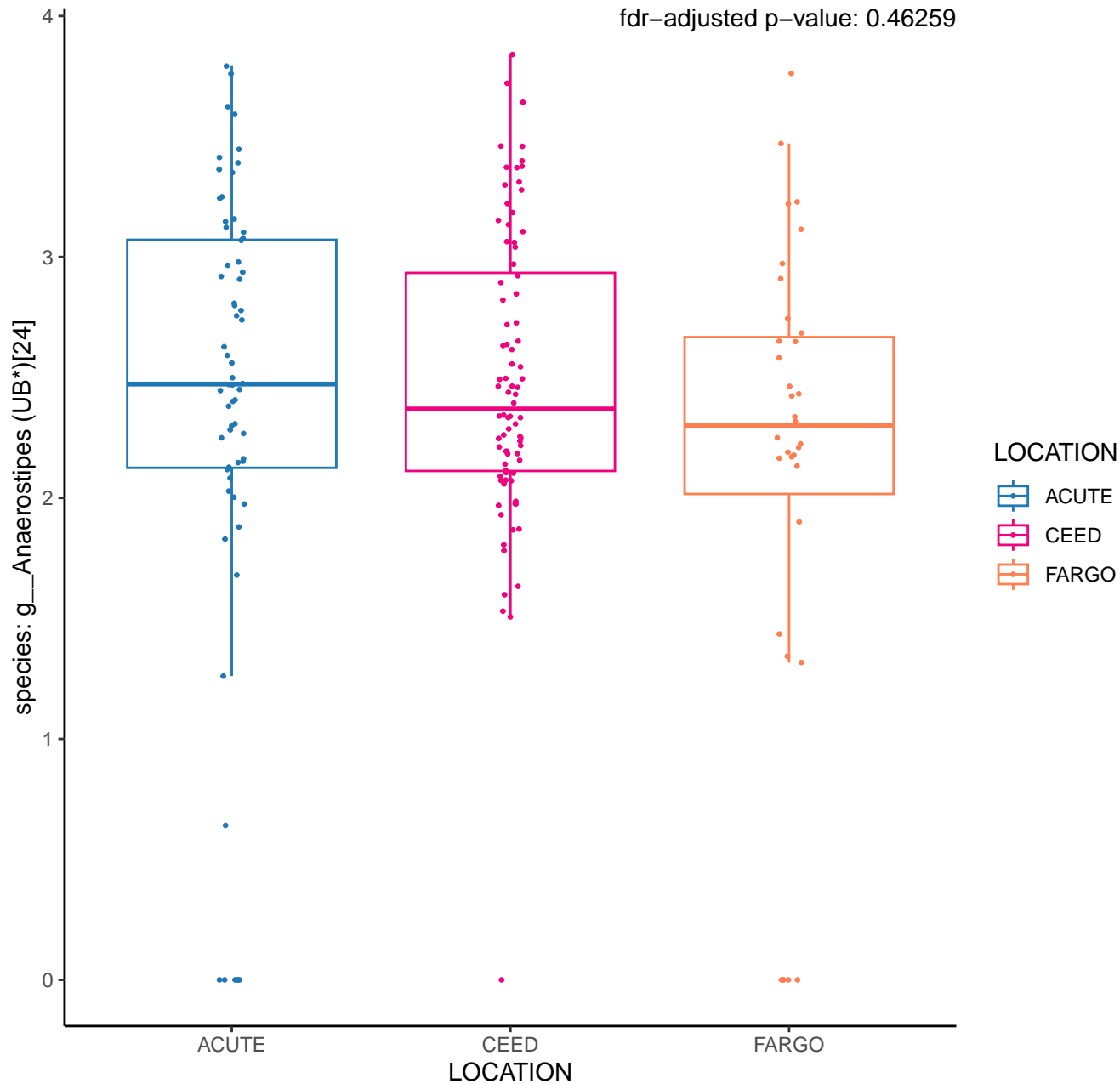
CEED

FARGO



anova p-value: 0.077417

fdr-adjusted p-value: 0.46259



anova p-value: 0.089879

fdr-adjusted p-value: 0.46259

species: Eubacterium.[38]

LOCATION

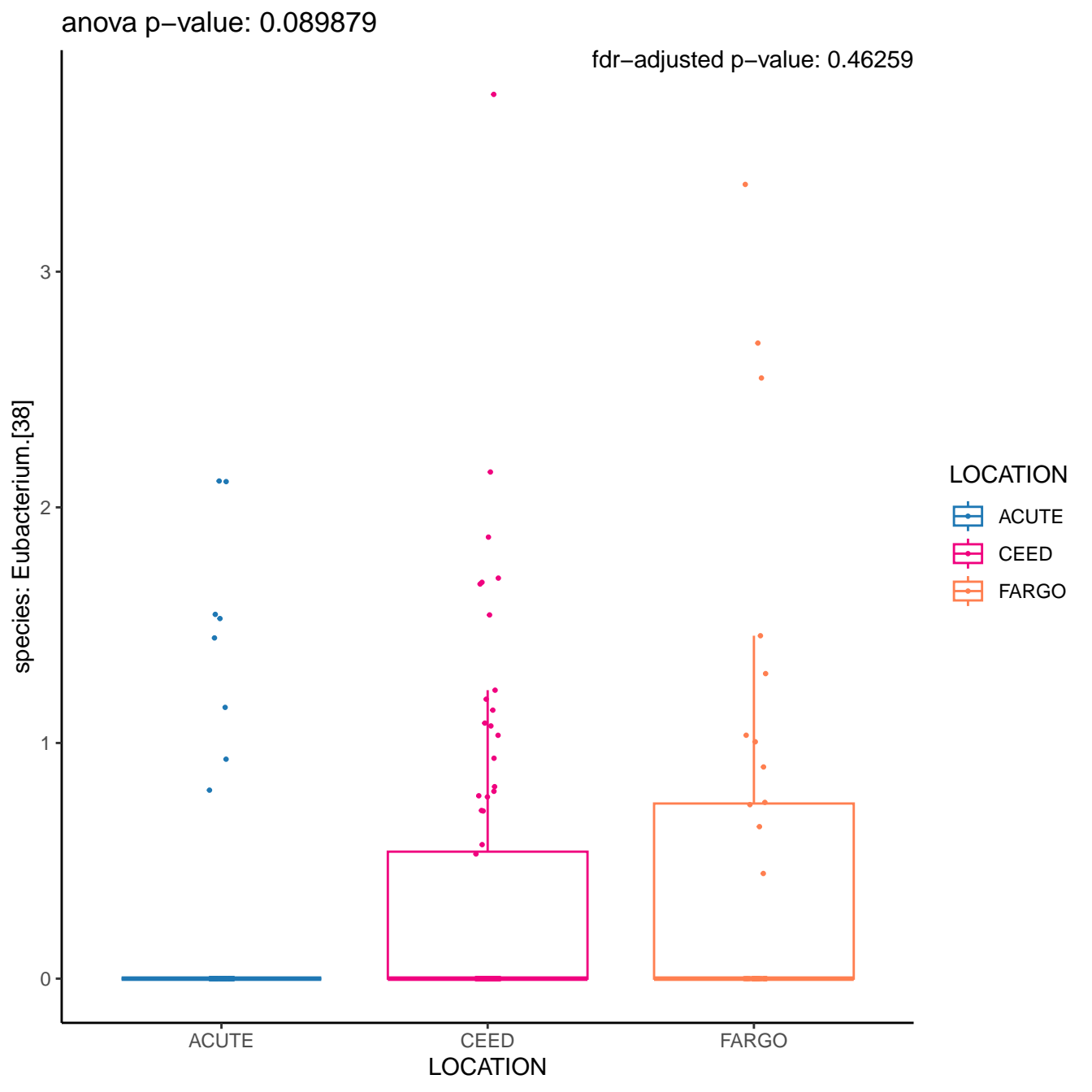


ACUTE

CEED

FARGO

LOCATION



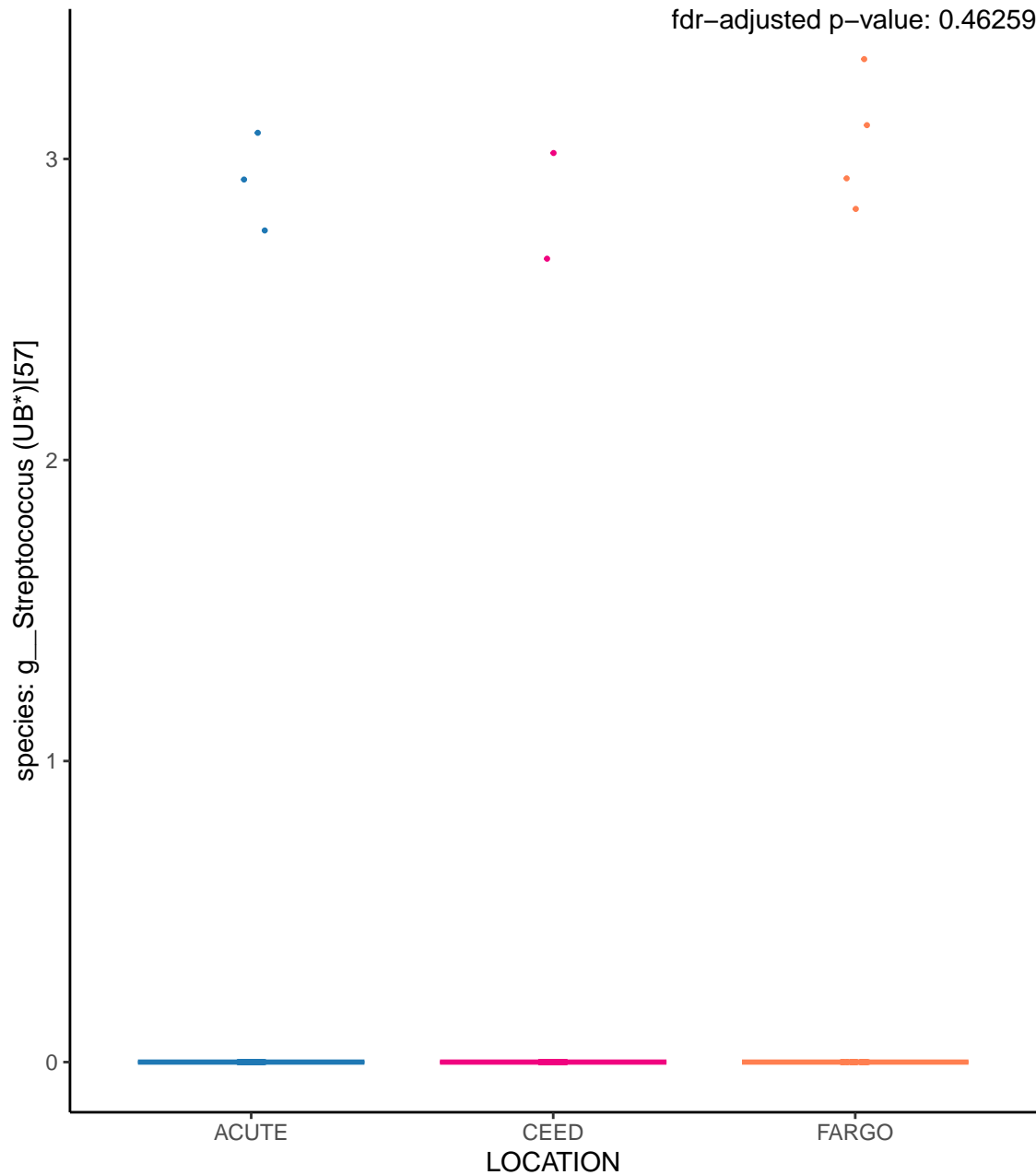
anova p-value: 0.093696

fdr-adjusted p-value: 0.46259

species: g__Streptococcus (UB*)[57]

LOCATION

ACUTE
CEED
FARGO



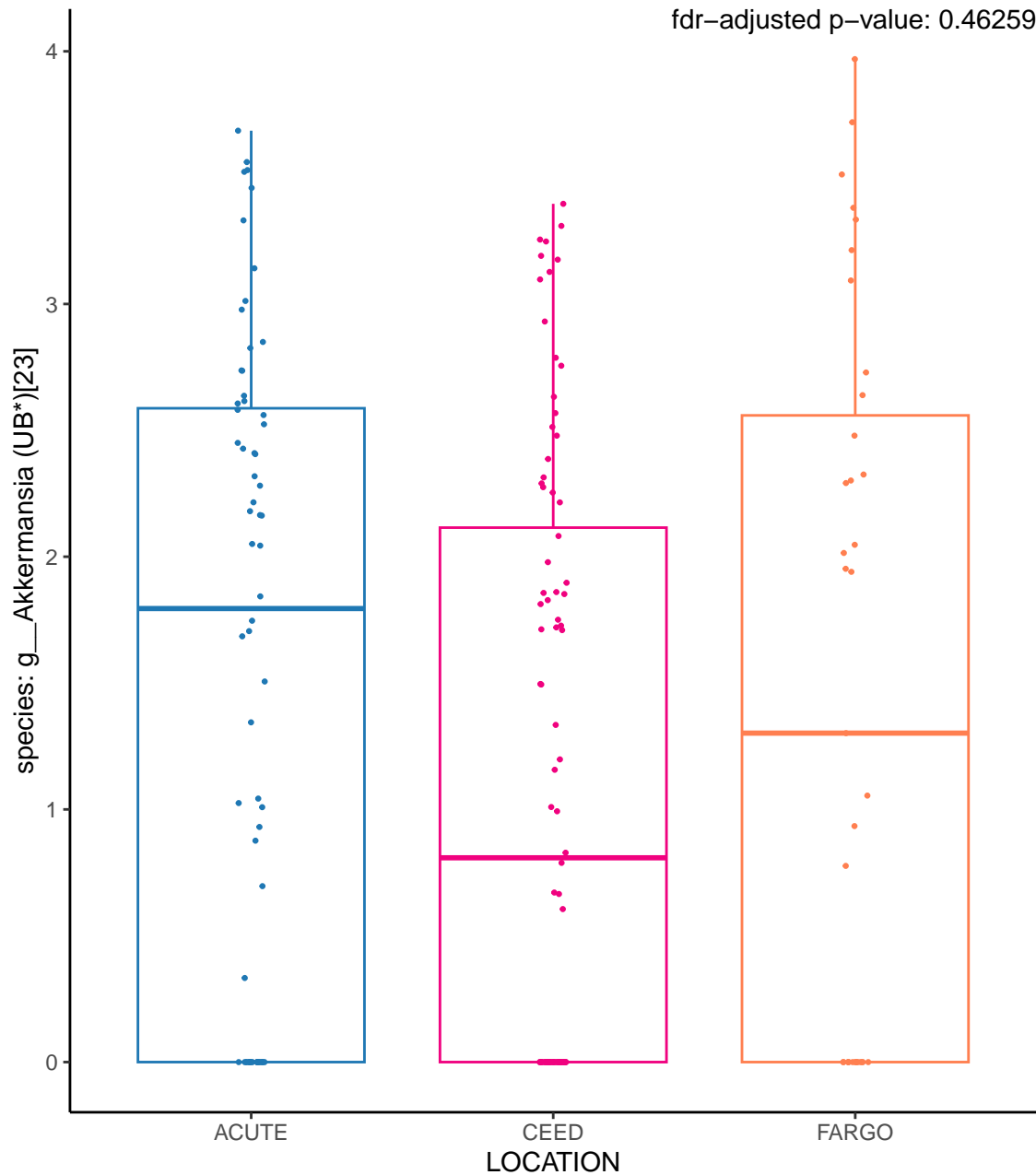
anova p-value: 0.094417

fdr-adjusted p-value: 0.46259

species: g__Akkermansia (UB*)[23]

LOCATION

ACUTE
CEED
FARGO



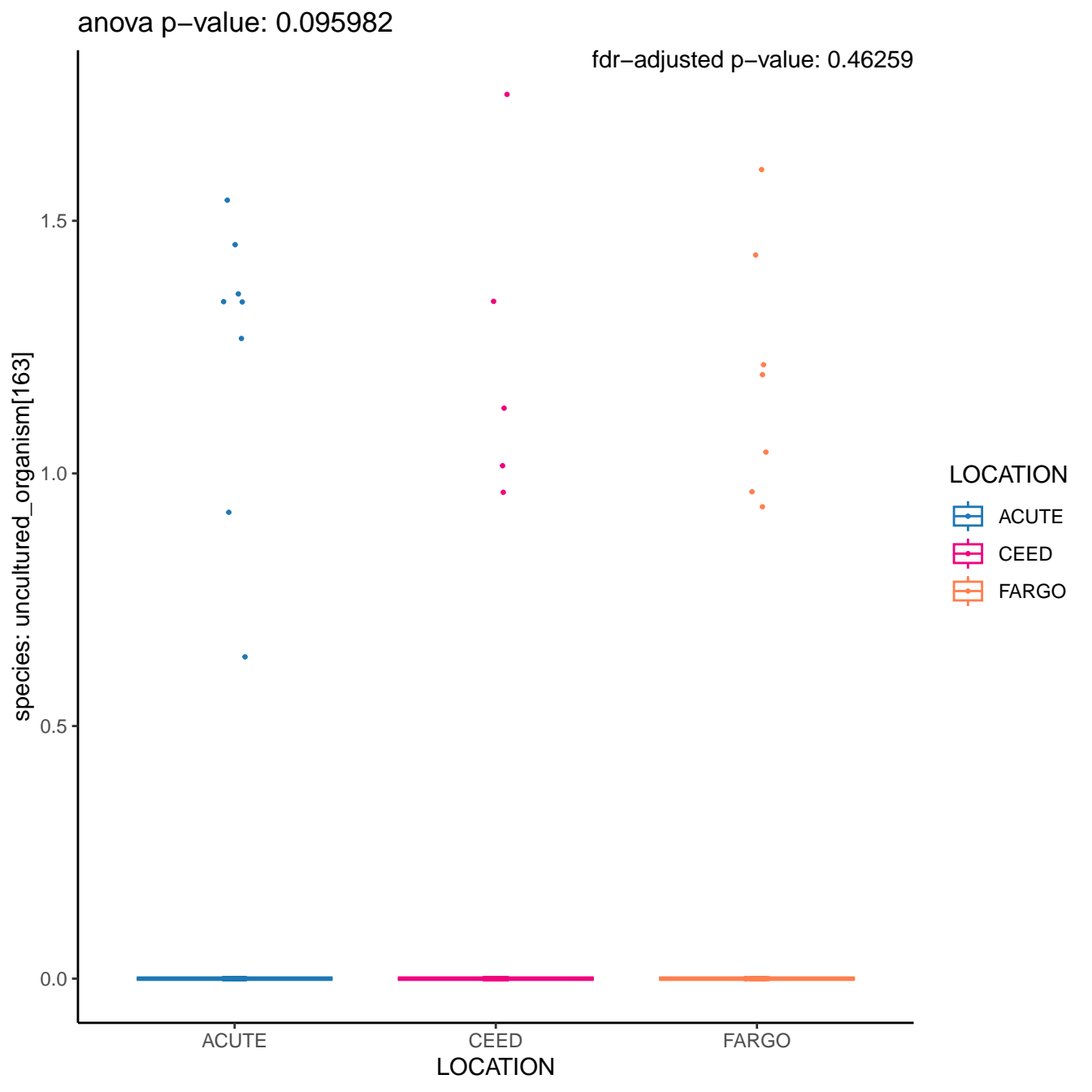
anova p-value: 0.095982

fdr-adjusted p-value: 0.46259

species: uncultured_organism[163]

LOCATION

ACUTE
CEED
FARGO



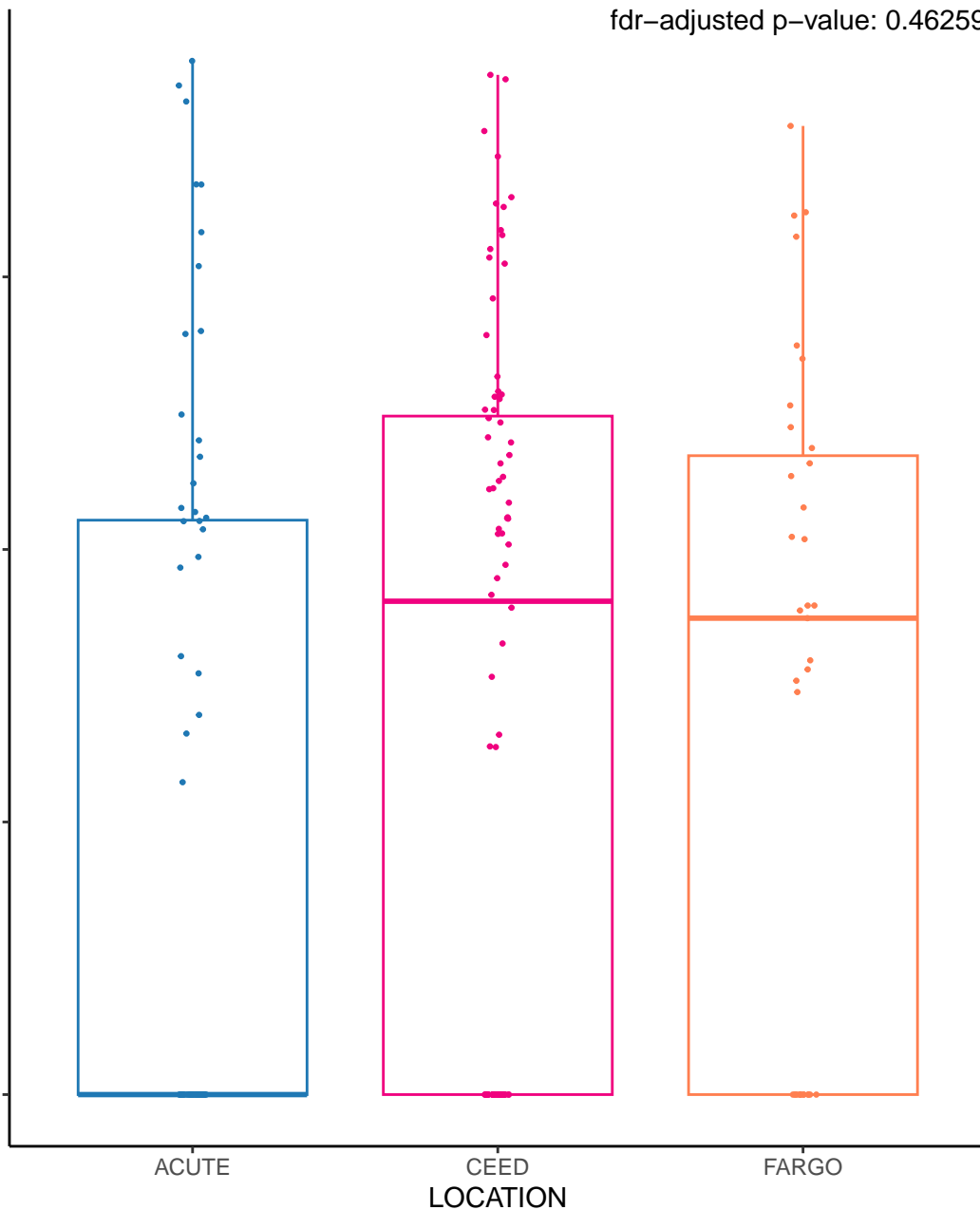
anova p-value: 0.096624

fdr-adjusted p-value: 0.46259

species: uncultured_organism[148]

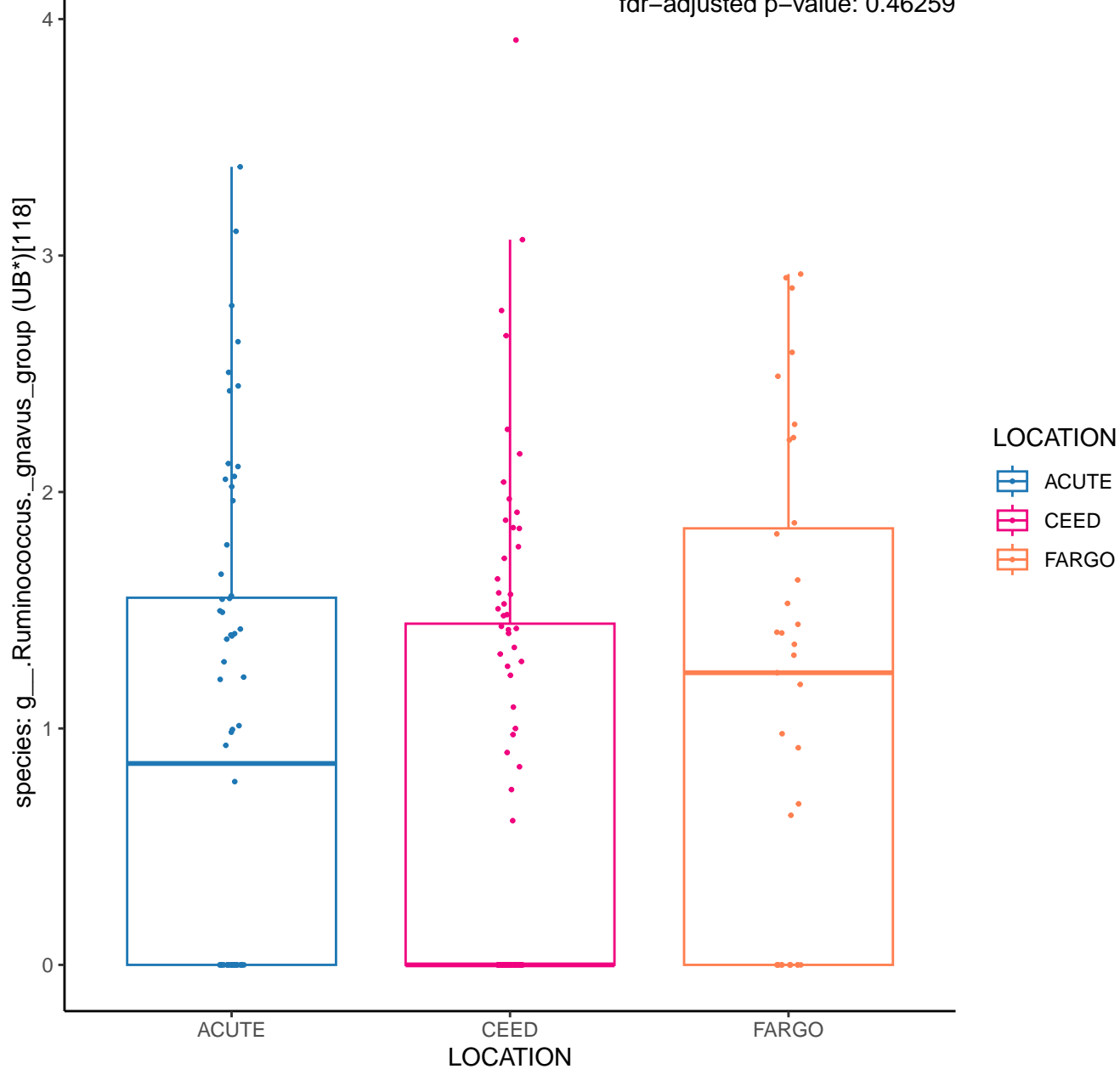
LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.098219

fdr-adjusted p-value: 0.46259

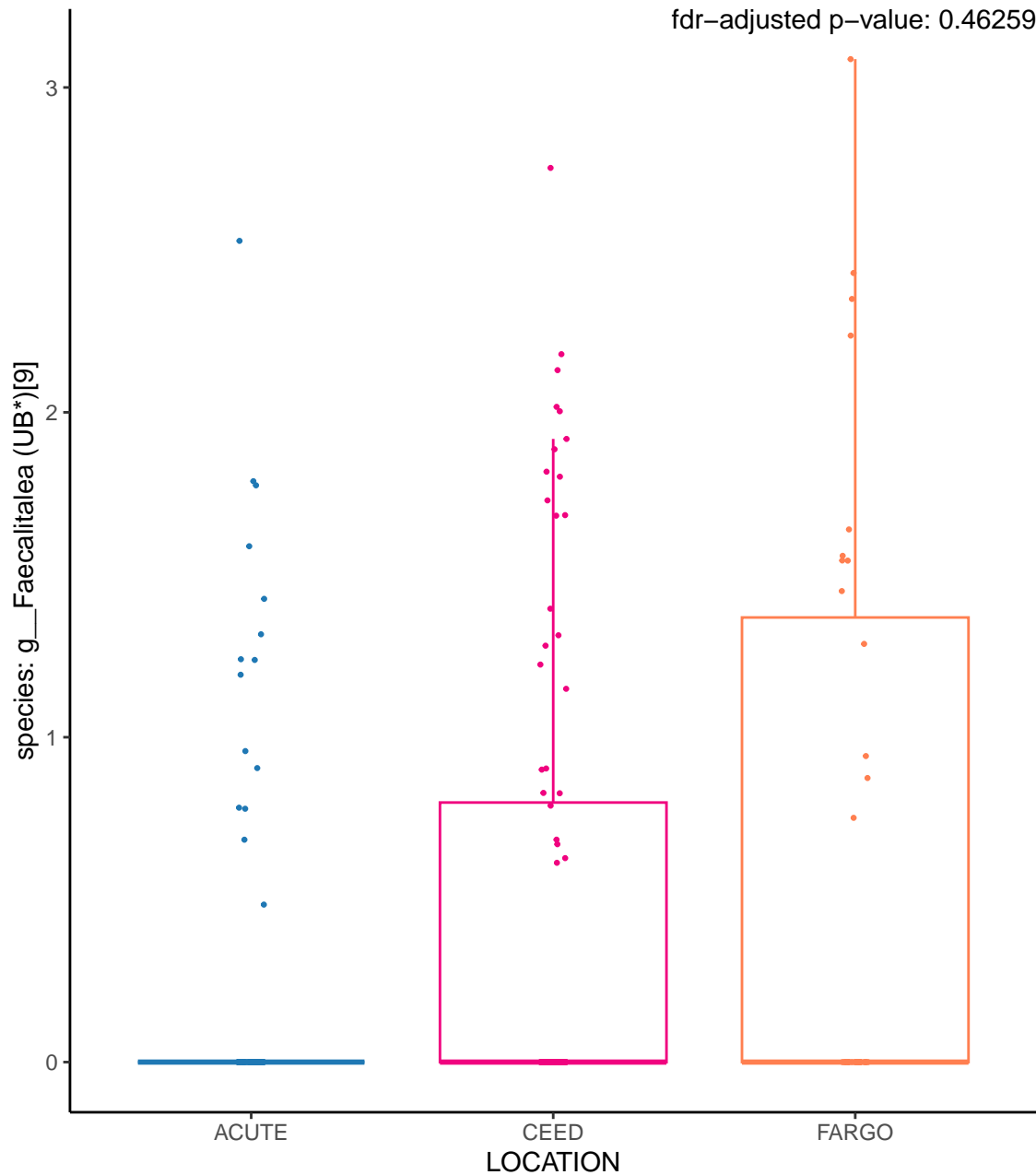


anova p-value: 0.098449

fdr-adjusted p-value: 0.46259

species: g__Faecalitalea (UB*)[9]

LOCATION



anova p-value: 0.10131

fdr-adjusted p-value: 0.46259

species: *Citrobacter*. [44]

LOCATION

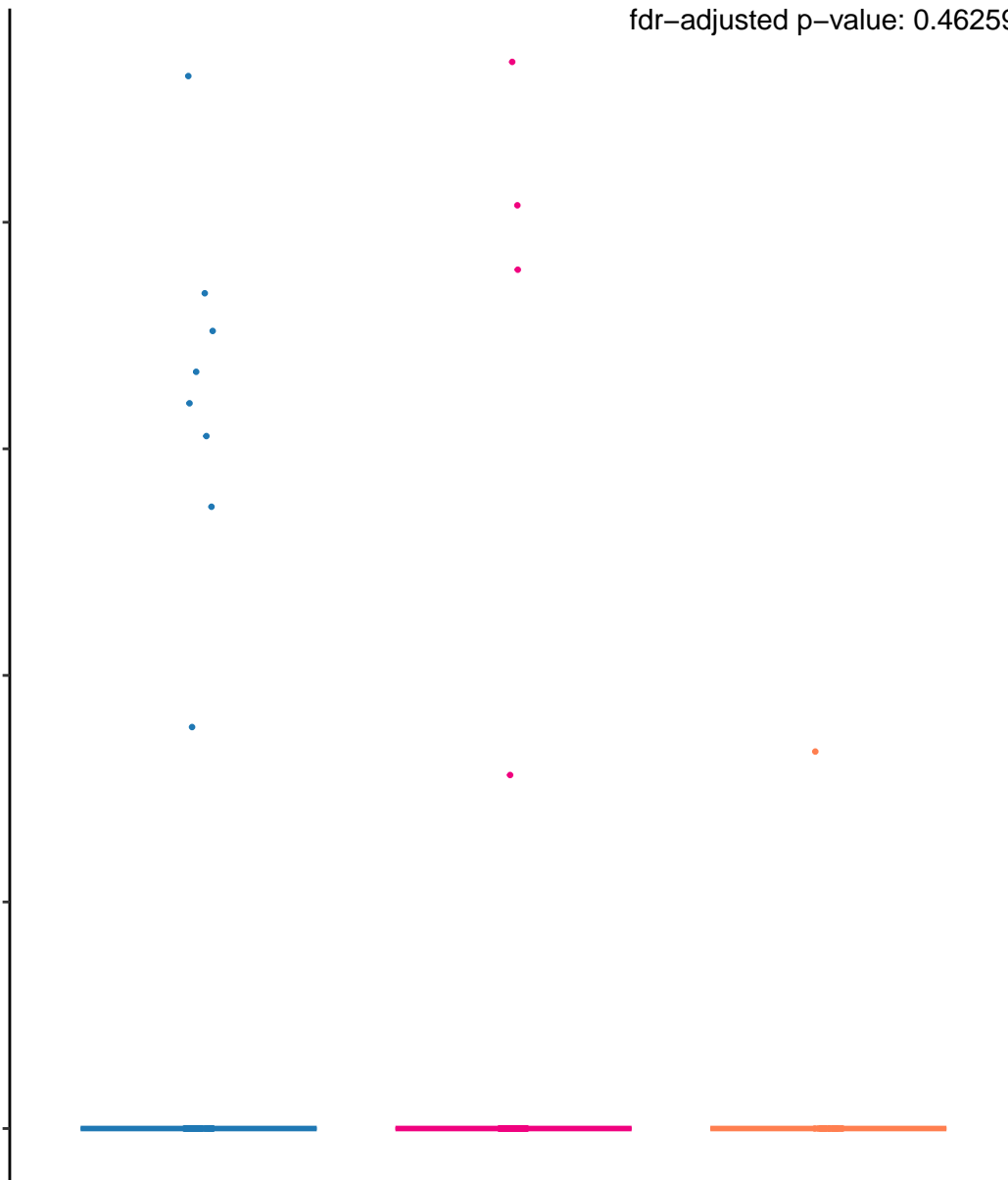
ACUTE
CEED
FARGO

ACUTE

CEED

FARGO

LOCATION



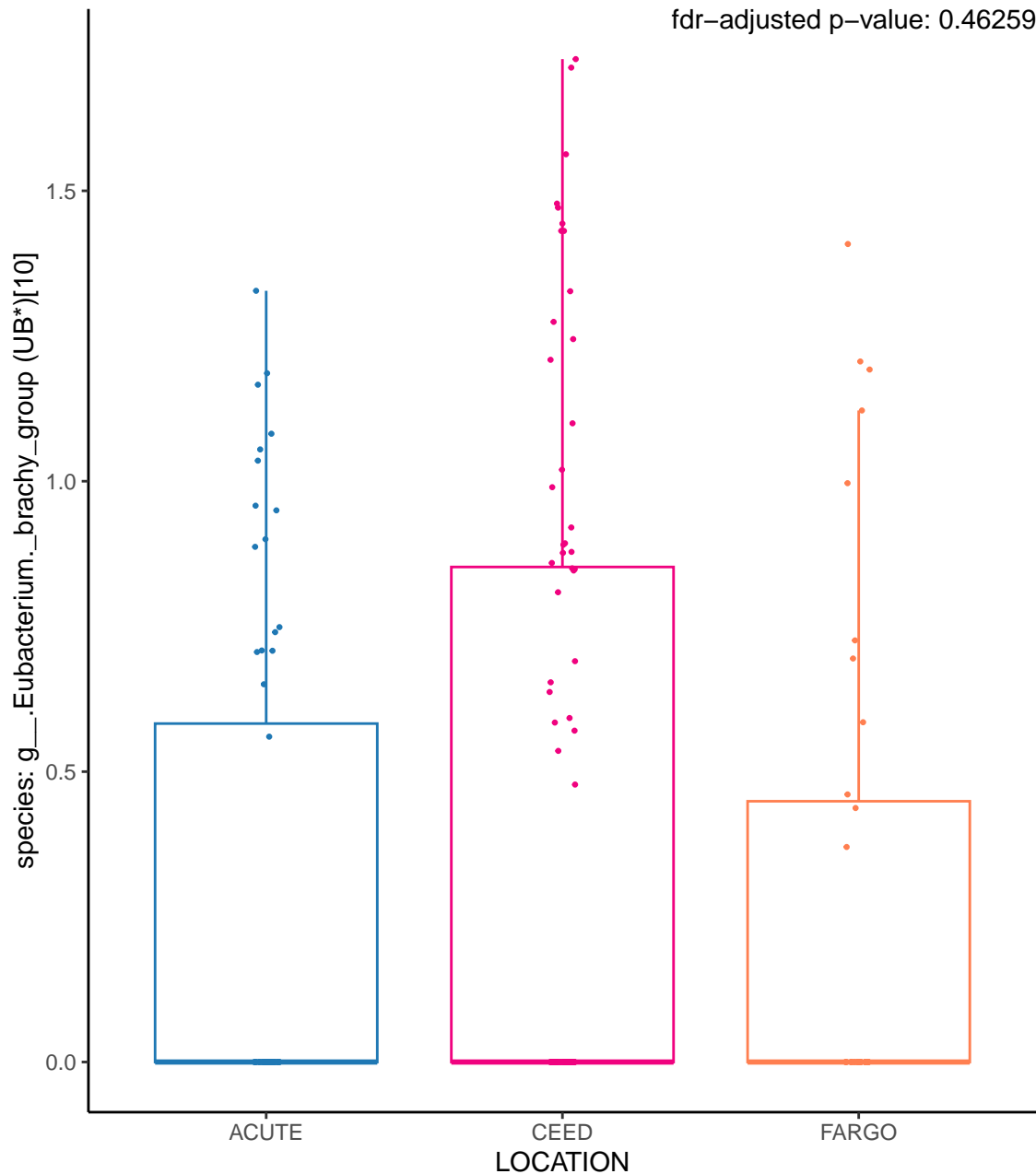
anova p-value: 0.10225

fdr-adjusted p-value: 0.46259

species: g__Eubacterium._brachy_group (UB*)[10]

LOCATION

ACUTE
CEED
FARGO



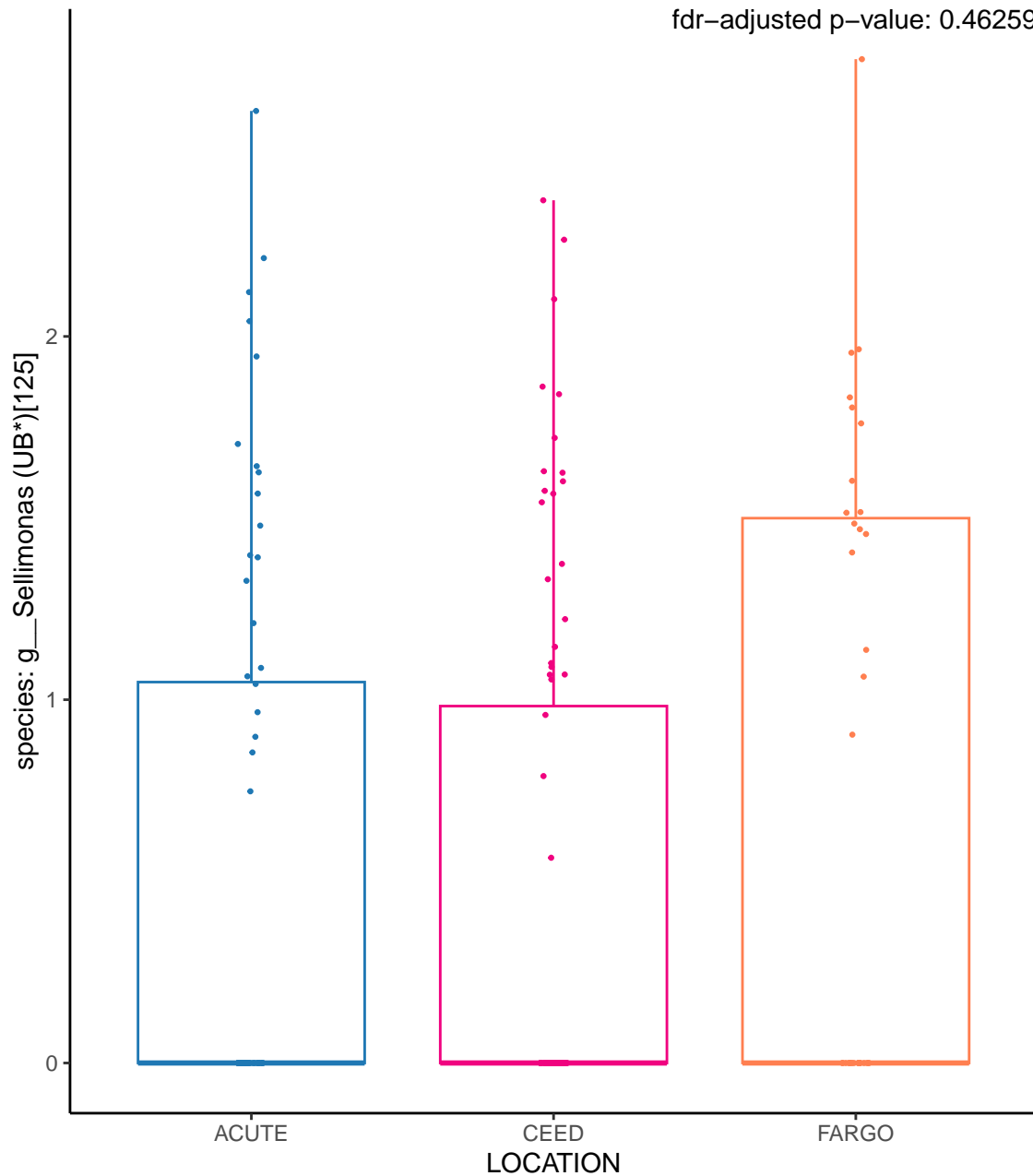
anova p-value: 0.10326

fdr-adjusted p-value: 0.46259

species: g__ Sellimonas (UB*)[125]

LOCATION

ACUTE
CEED
FARGO

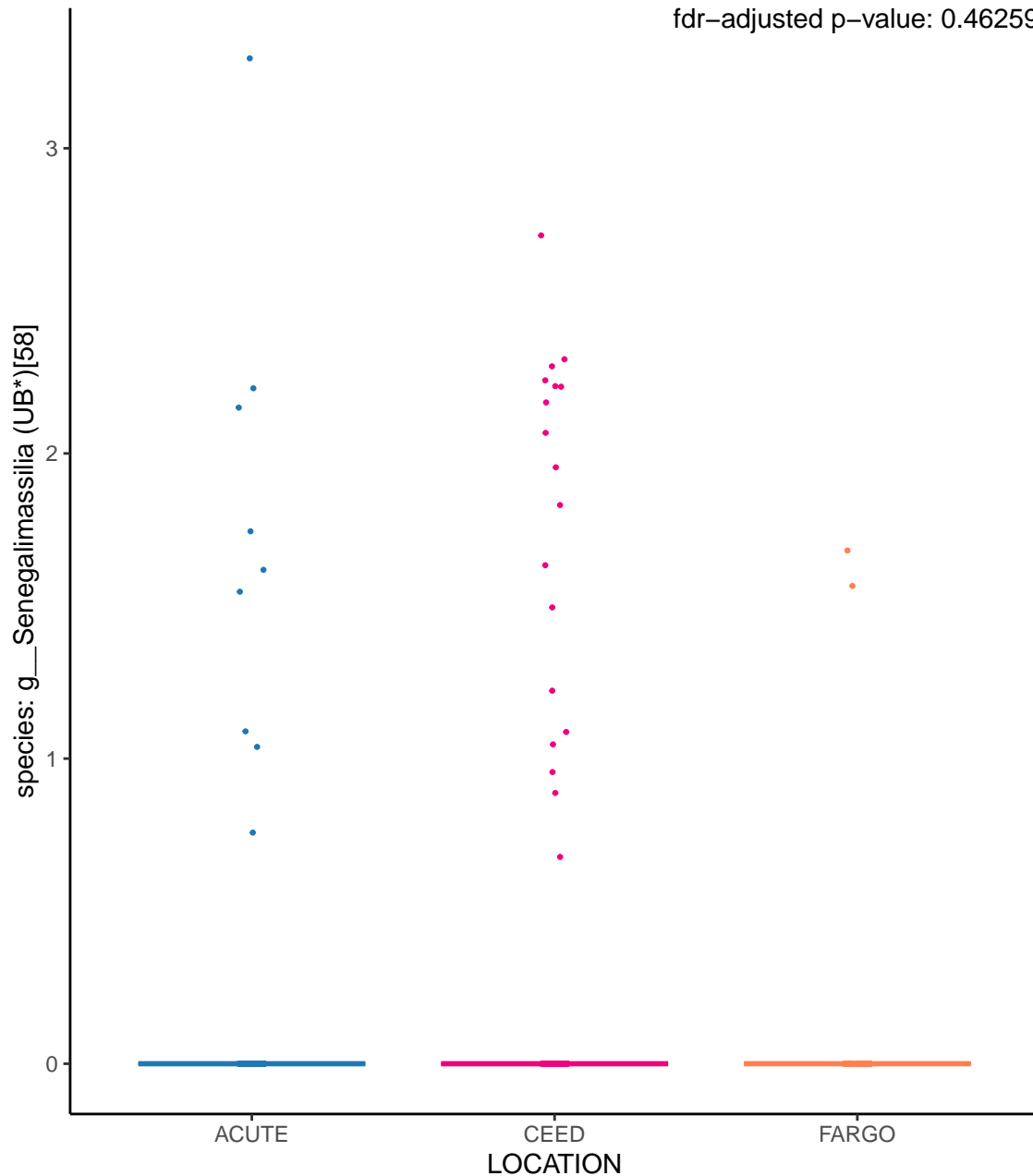
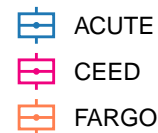


anova p-value: 0.1136

fdr-adjusted p-value: 0.46259

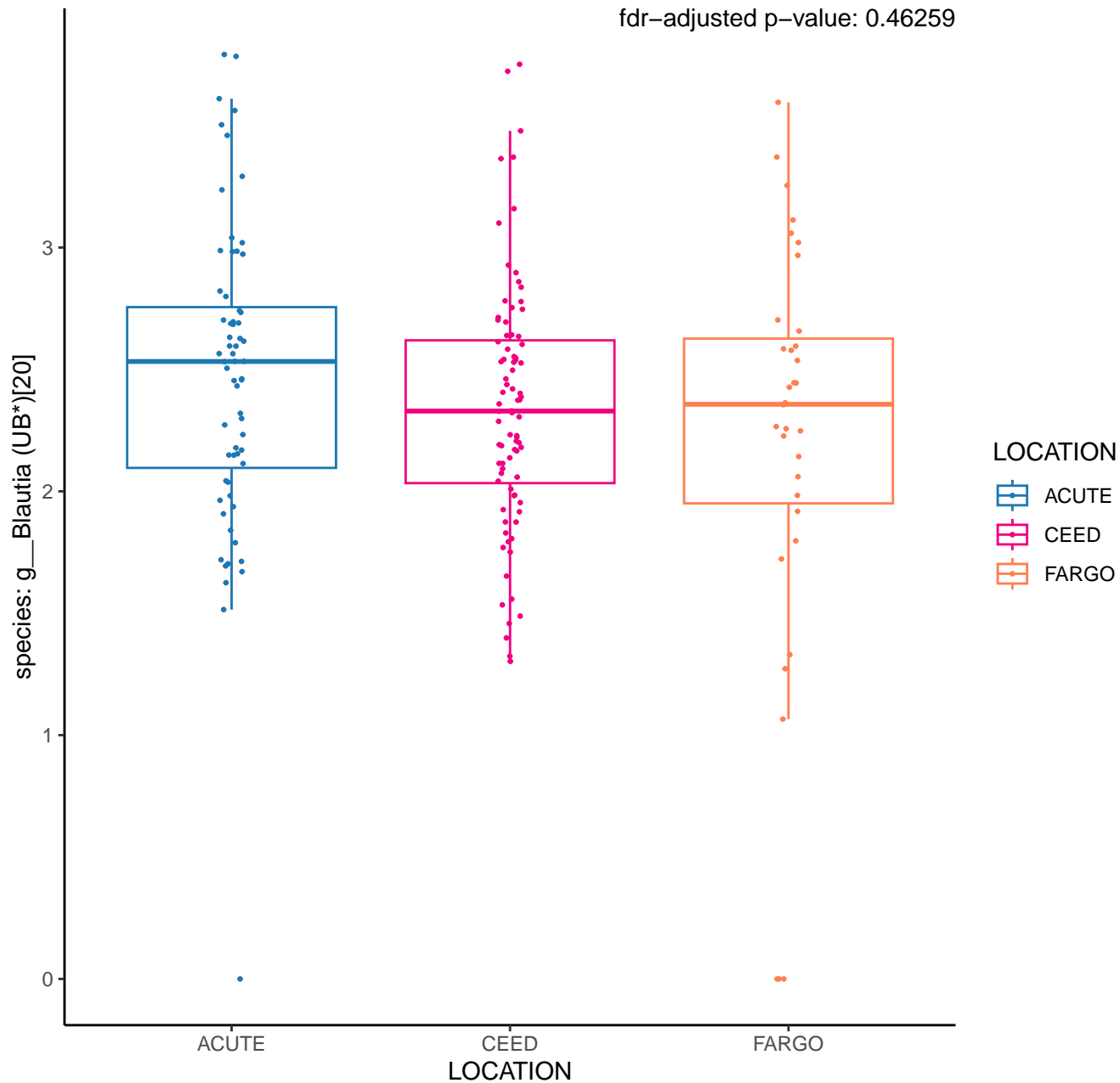
species: g__Senegalimassilia (UB*)[58]

LOCATION



anova p-value: 0.11427

fdr-adjusted p-value: 0.46259



anova p-value: 0.11456

fdr-adjusted p-value: 0.46259

species: g__Erysipelatoclostridium (UB*)[31]

LOCATION

ACUTE

CEED

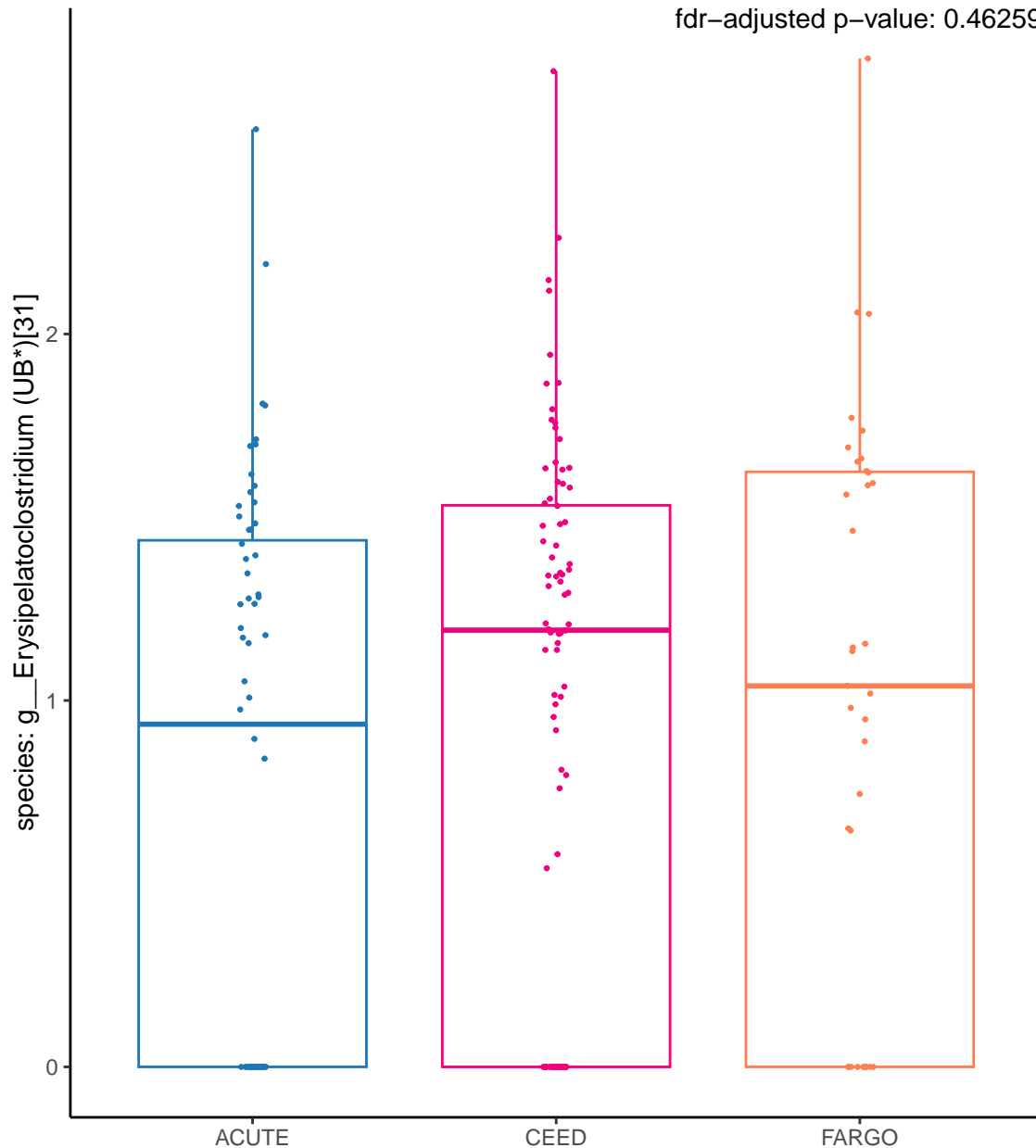
FARGO

ACUTE

CEED

FARGO

LOCATION



anova p-value: 0.12055

fdr-adjusted p-value: 0.46259

species: g__Escherichia.Shigella (UB*)[174]

LOCATION

ACUTE

CEED

FARGO

0

ACUTE

CEED

FARGO

LOCATION

3

2

1

anova p-value: 0.12055

fdr-adjusted p-value: 0.46259

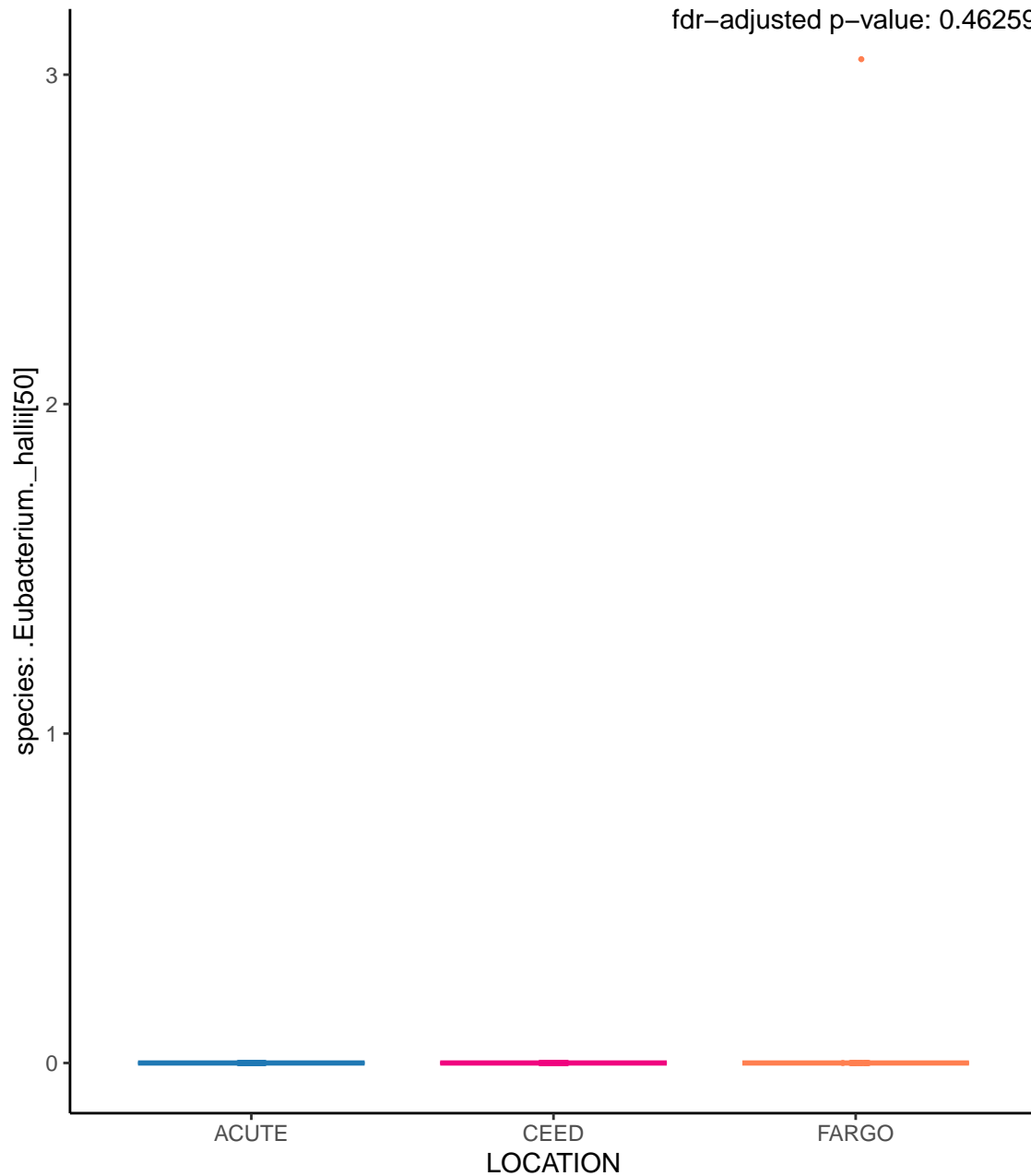
species: .Eubacterium._hallii[50]

LOCATION

ACUTE

CEED

FARGO



anova p-value: 0.12055

fdr-adjusted p-value: 0.46259

species: Bifidobacterium_longum[140]

LOCATION

ACUTE

CEED

FARGO

0

ACUTE

CEED

FARGO

LOCATION

3

2

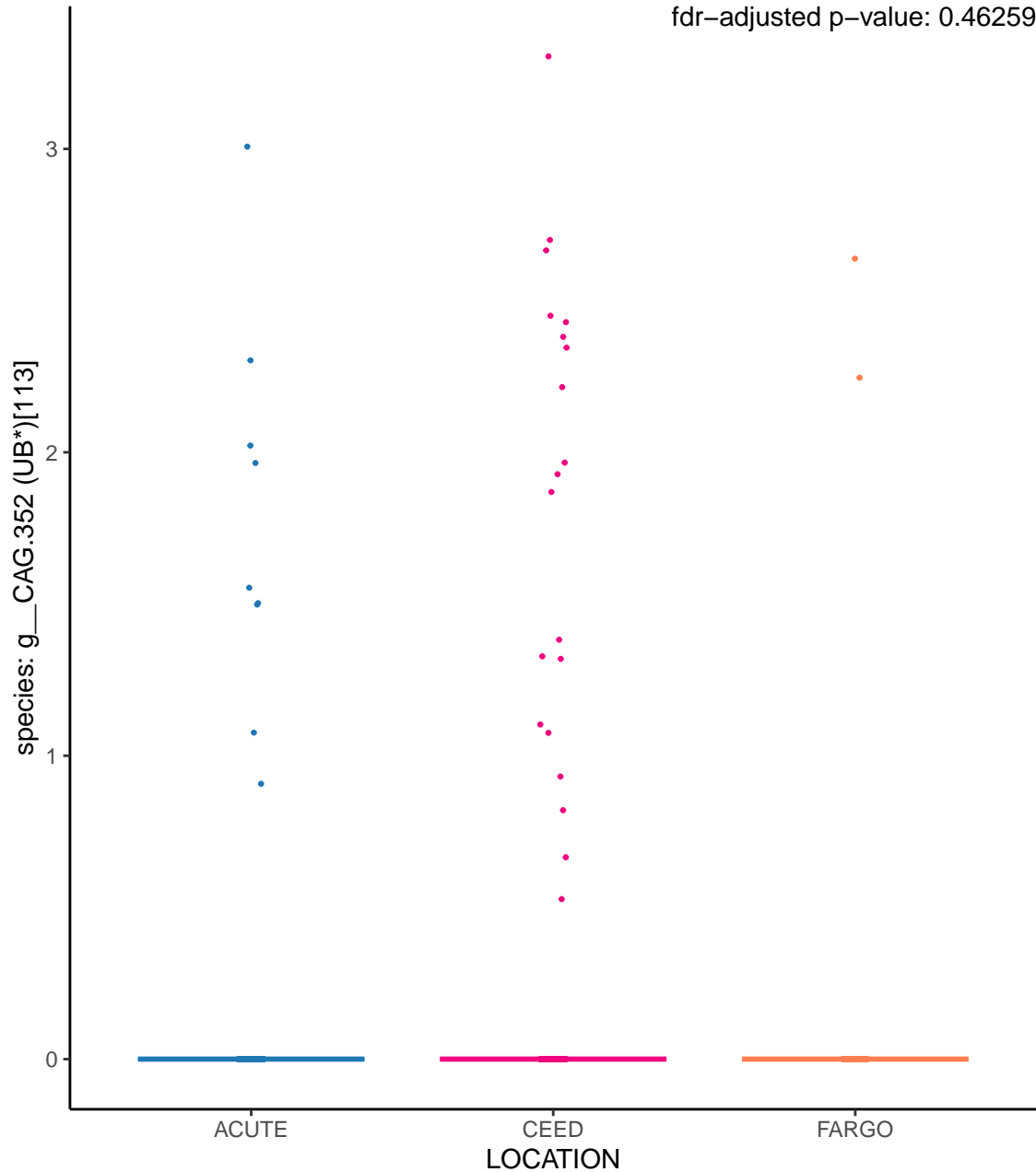
1

anova p-value: 0.12317

fdr-adjusted p-value: 0.46259

species: g__CAG.352 (UB*)[113]

LOCATION



anova p-value: 0.12333

fdr-adjusted p-value: 0.46259

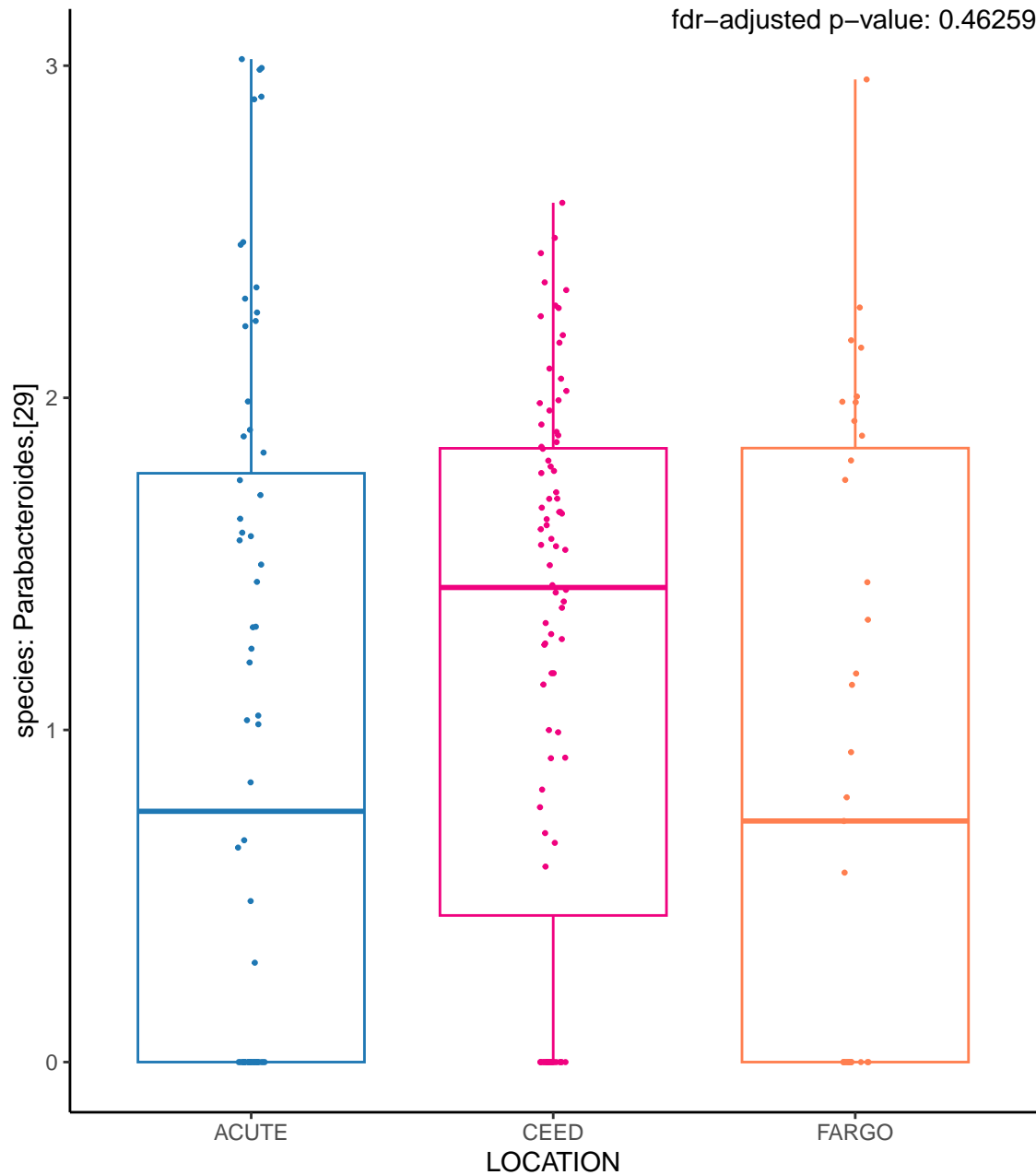
species: Parabacteroides.[29]

LOCATION

ACUTE

CEED

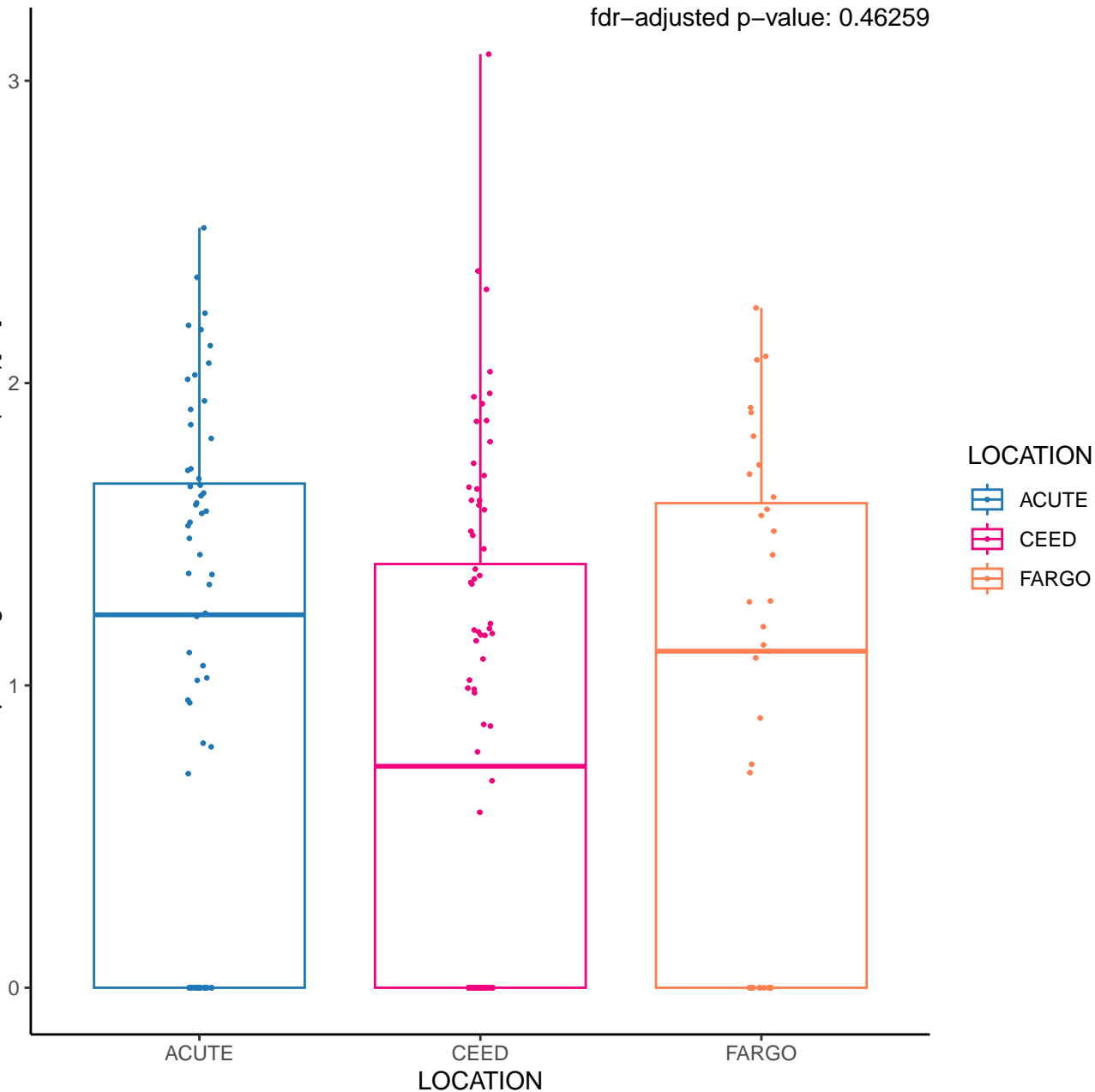
FARGO



anova p-value: 0.12445

fdr-adjusted p-value: 0.46259

species: g__Flavonifractor (UB*)[47]



anova p-value: 0.12805

fdr-adjusted p-value: 0.46259

species: g__Slackia (UB*)[133]

LOCATION

ACUTE

CEED

FARGO

0

1

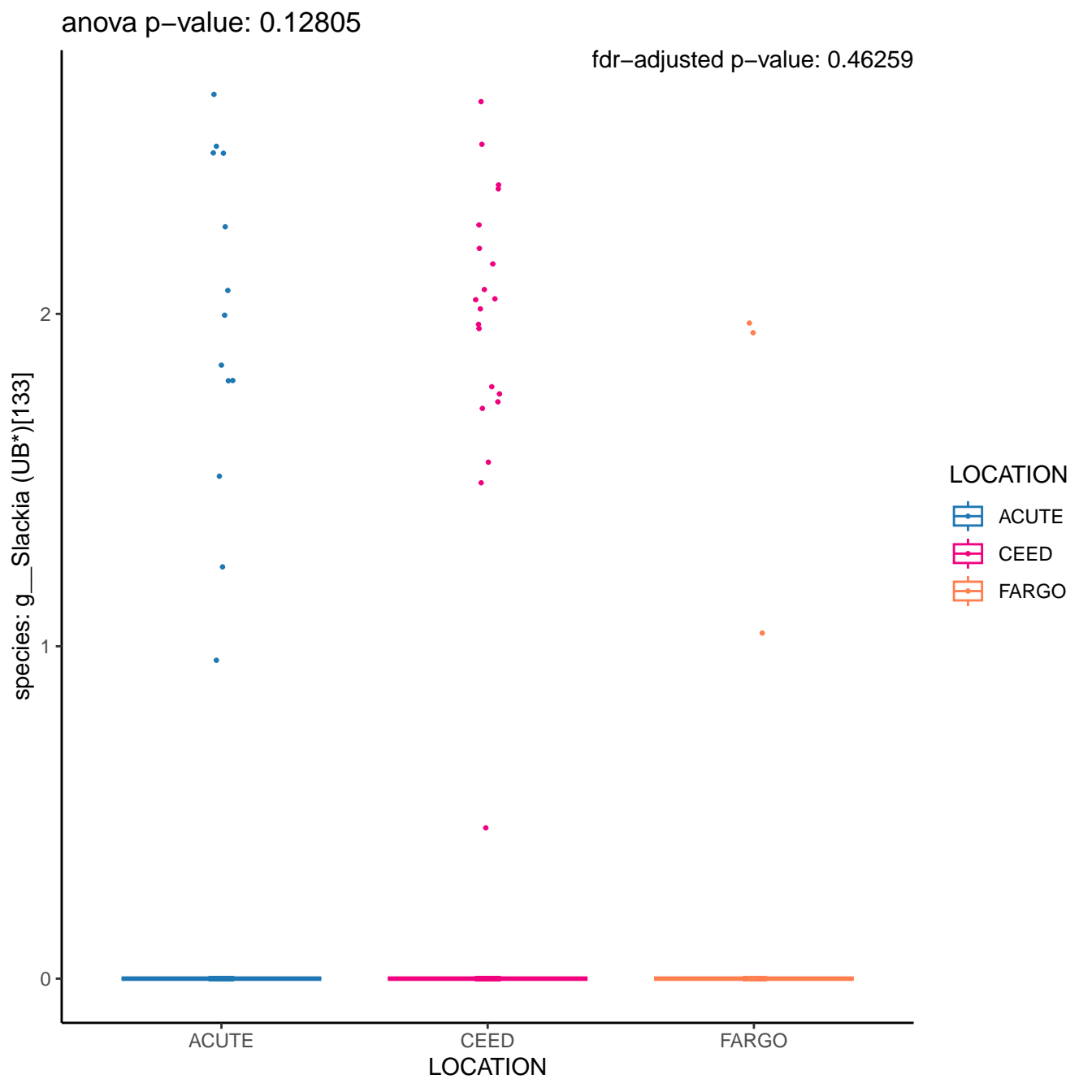
2

ACUTE

CEED

FARGO

LOCATION



anova p-value: 0.12985

fdr-adjusted p-value: 0.46259

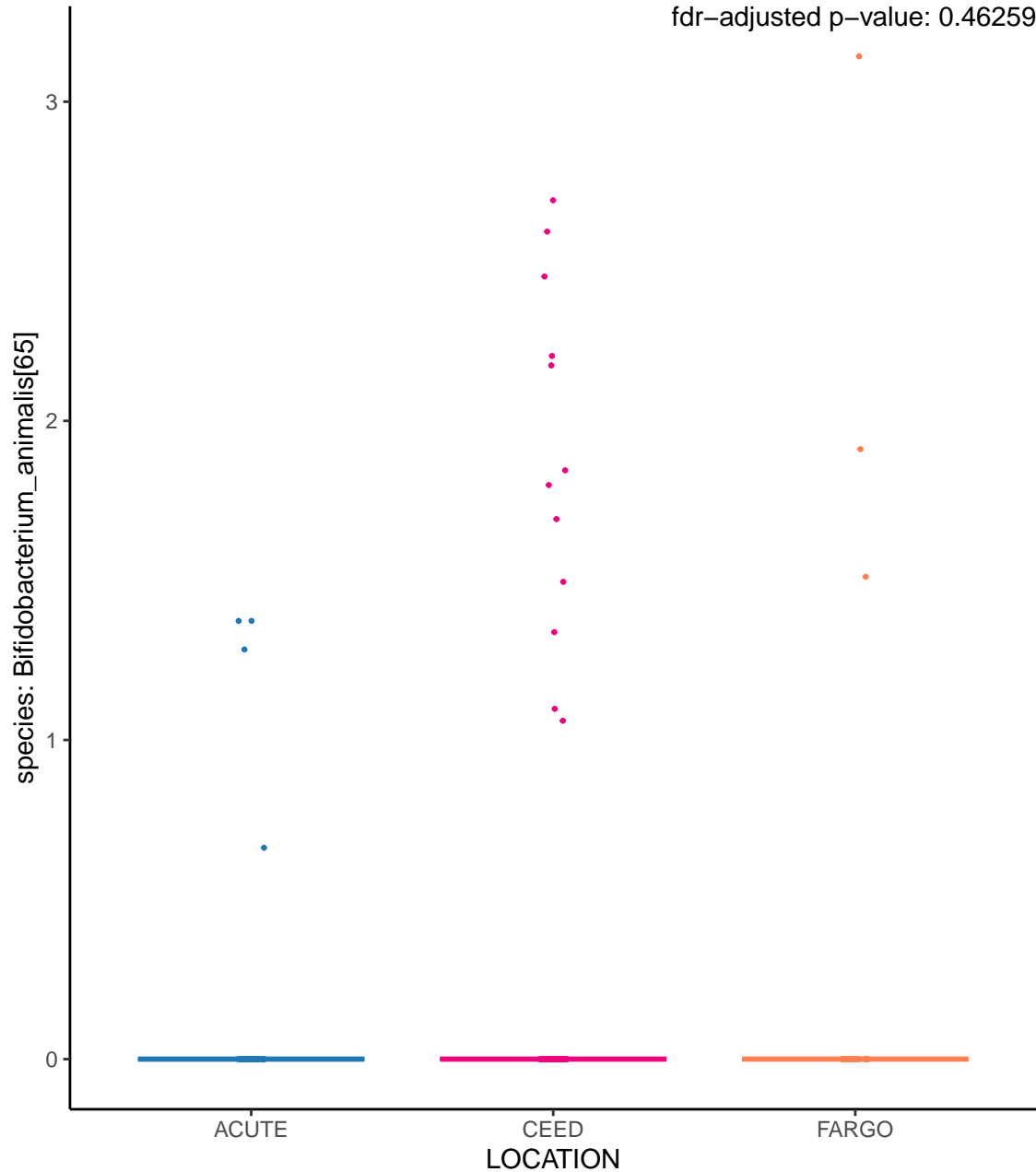
species: *Bifidobacterium animalis*[65]

LOCATION

ACUTE

CEED

FARGO



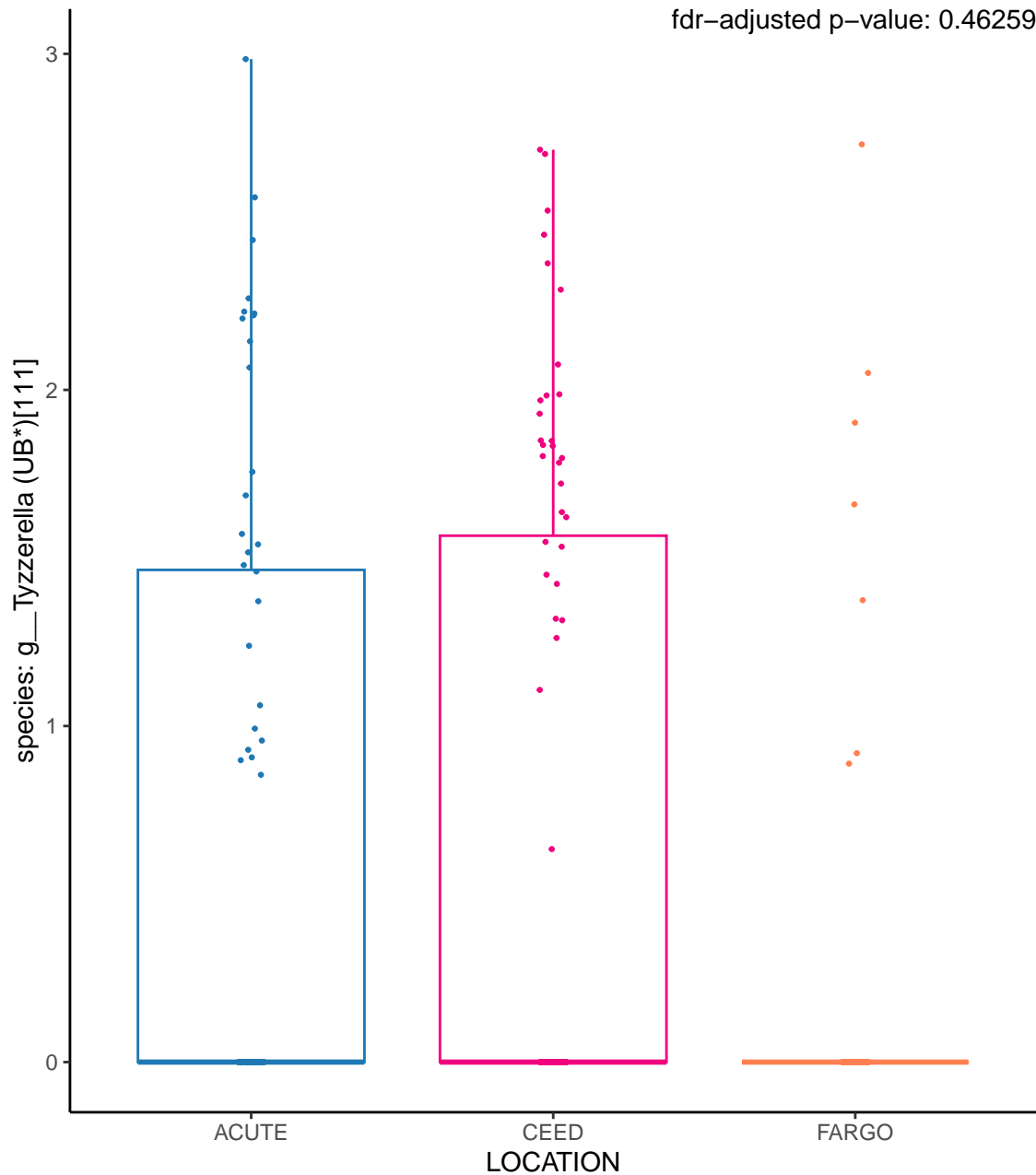
anova p-value: 0.13341

fdr-adjusted p-value: 0.46259

species: g__Tyzzerella (UB*)[111]

LOCATION

ACUTE
CEED
FARGO

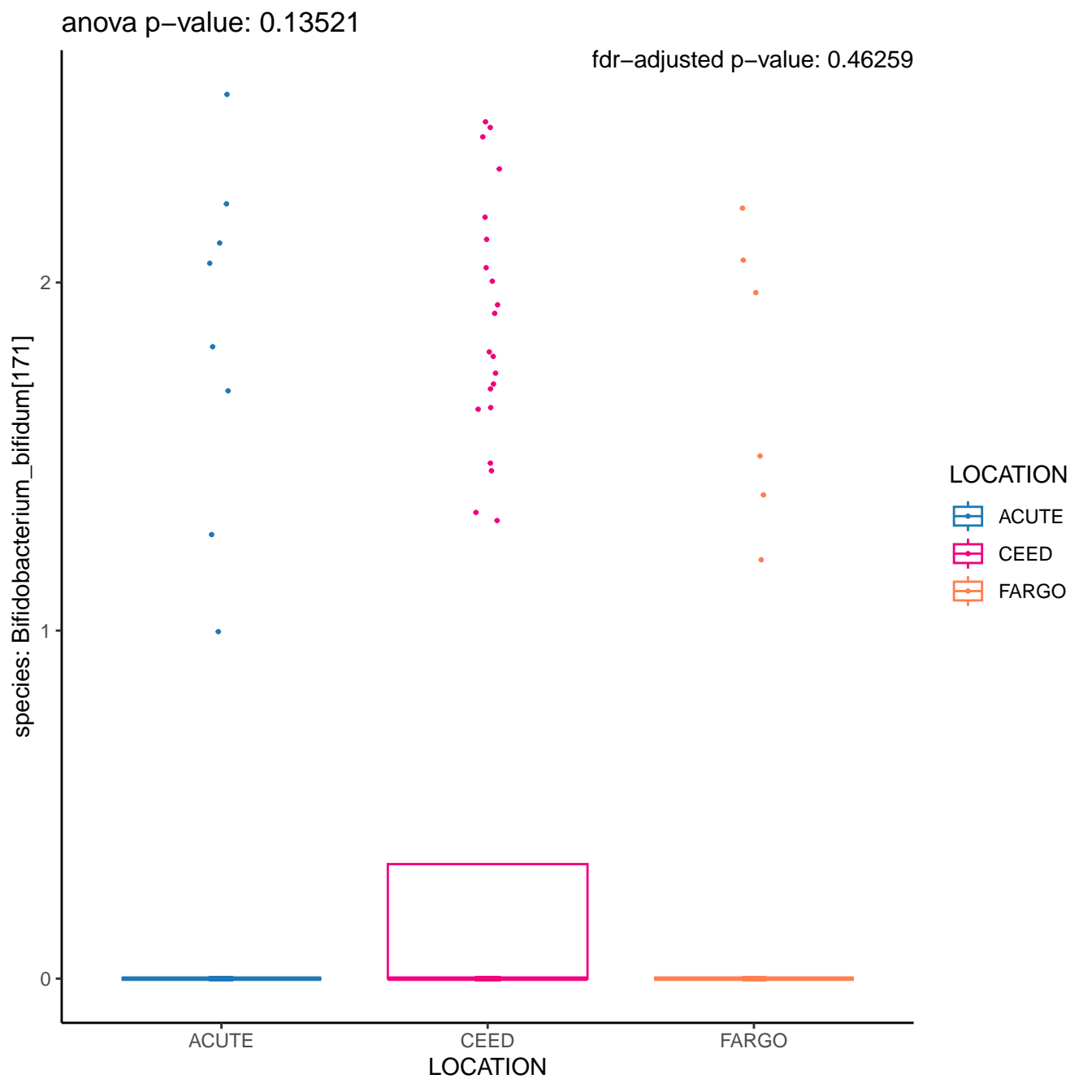
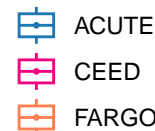


anova p-value: 0.13521

fdr-adjusted p-value: 0.46259

species: *Bifidobacterium_bifidum*[171]

LOCATION



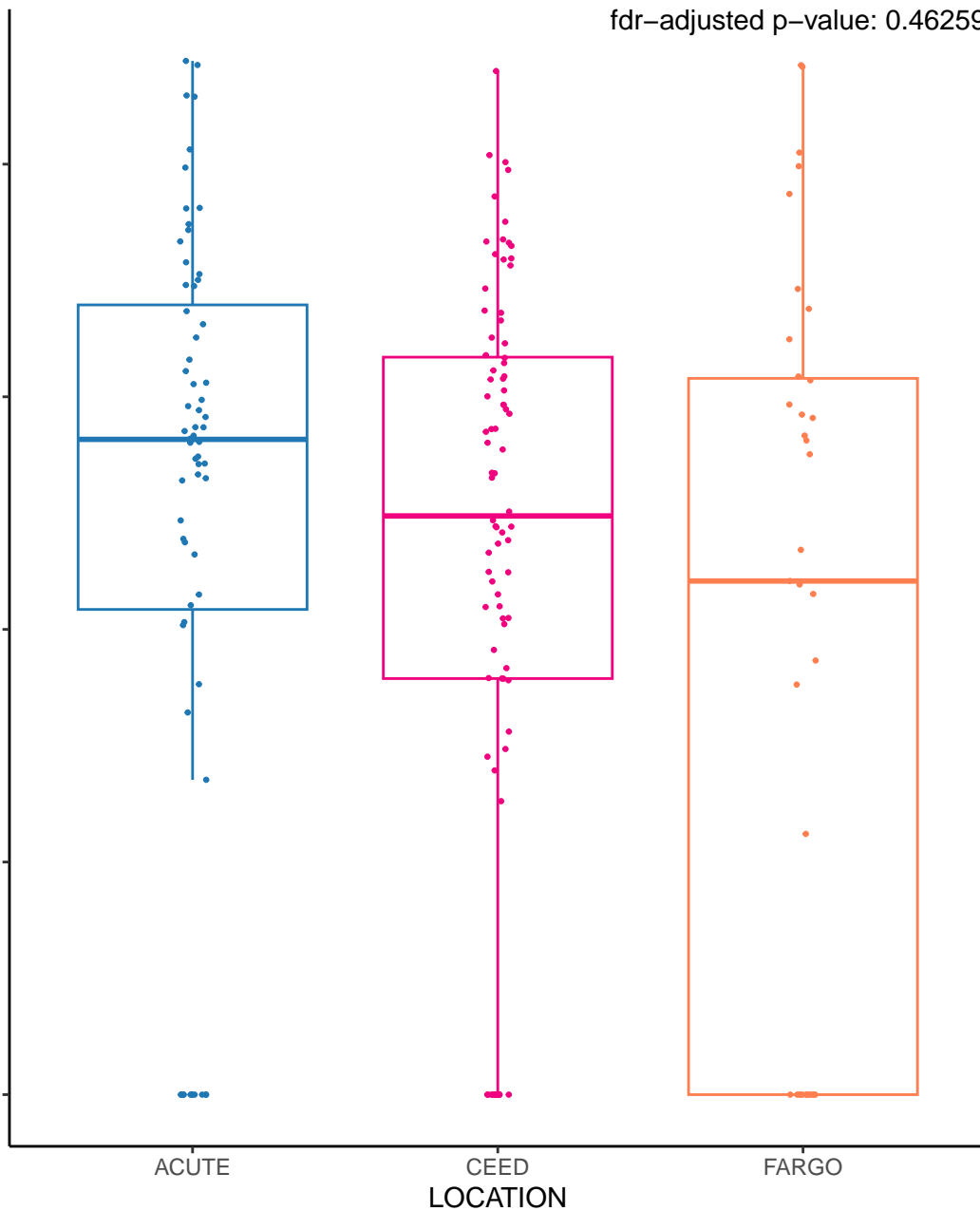
anova p-value: 0.13716

fdr-adjusted p-value: 0.46259

species: g_uncultured (UB*)[61]

LOCATION

ACUTE
CEED
FARGO



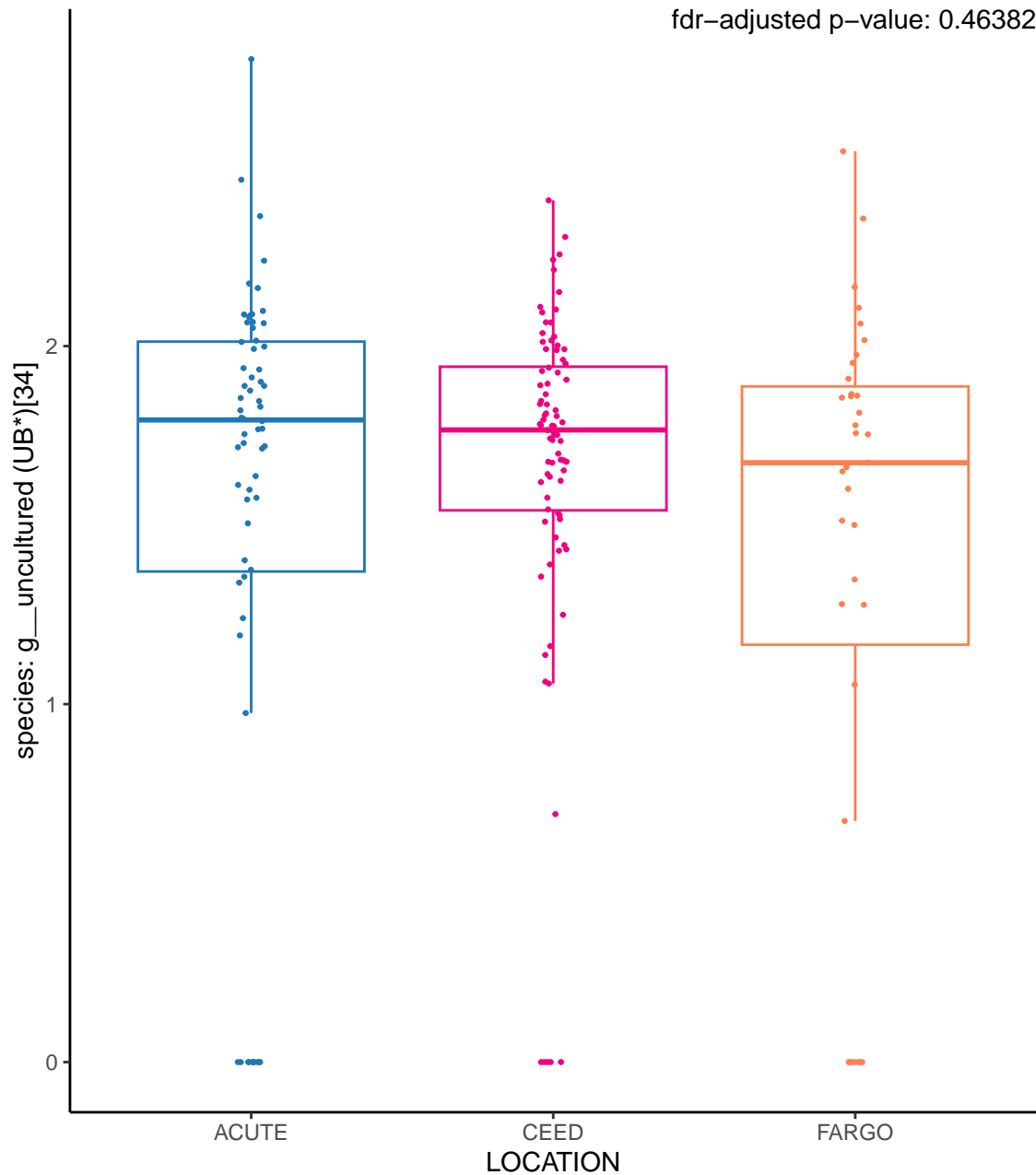
anova p-value: 0.14022

fdr-adjusted p-value: 0.46382

species: g__uncultured (UB*)[34]

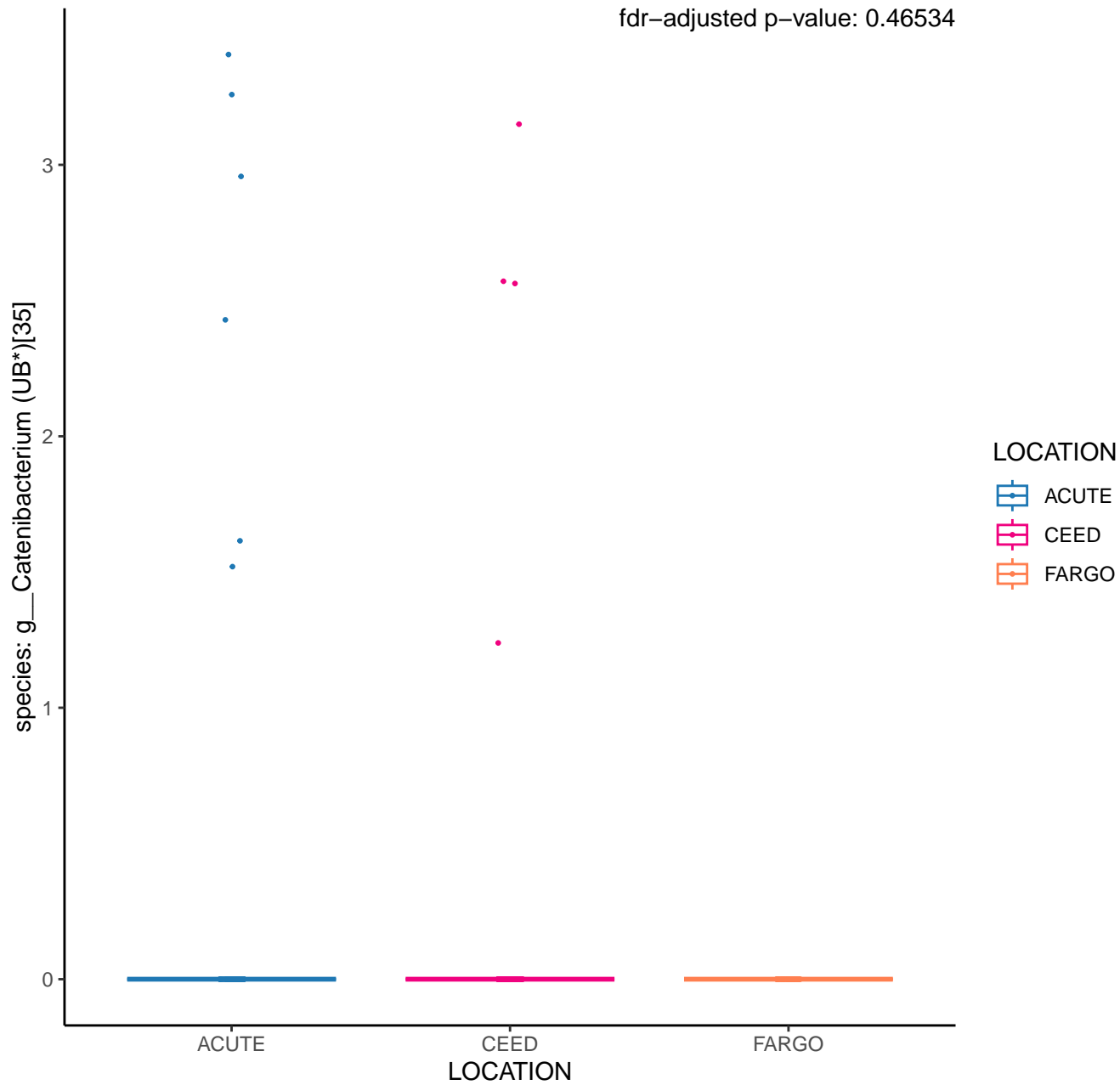
LOCATION

ACUTE
CEED
FARGO



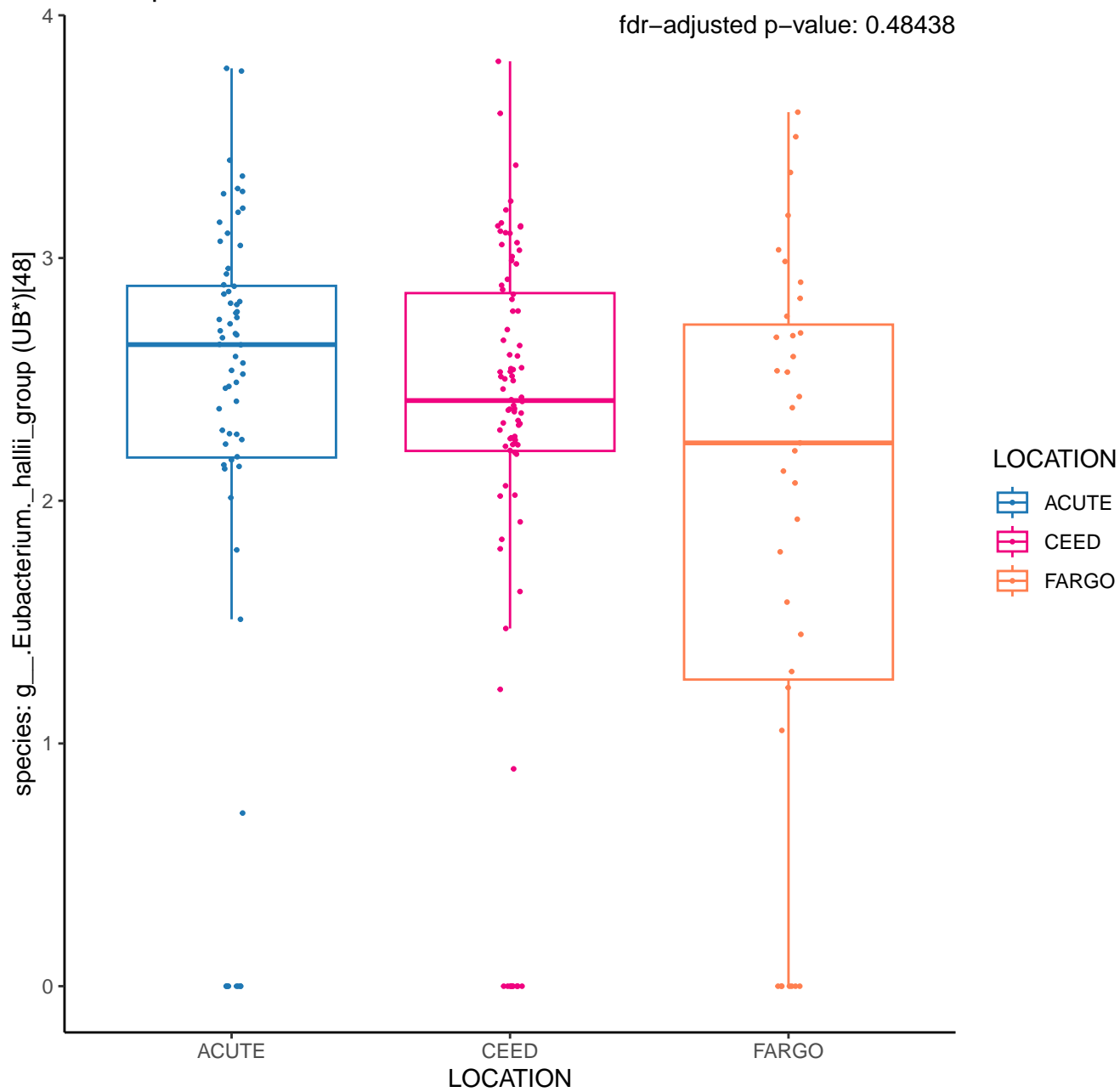
anova p-value: 0.14339

fdr-adjusted p-value: 0.46534



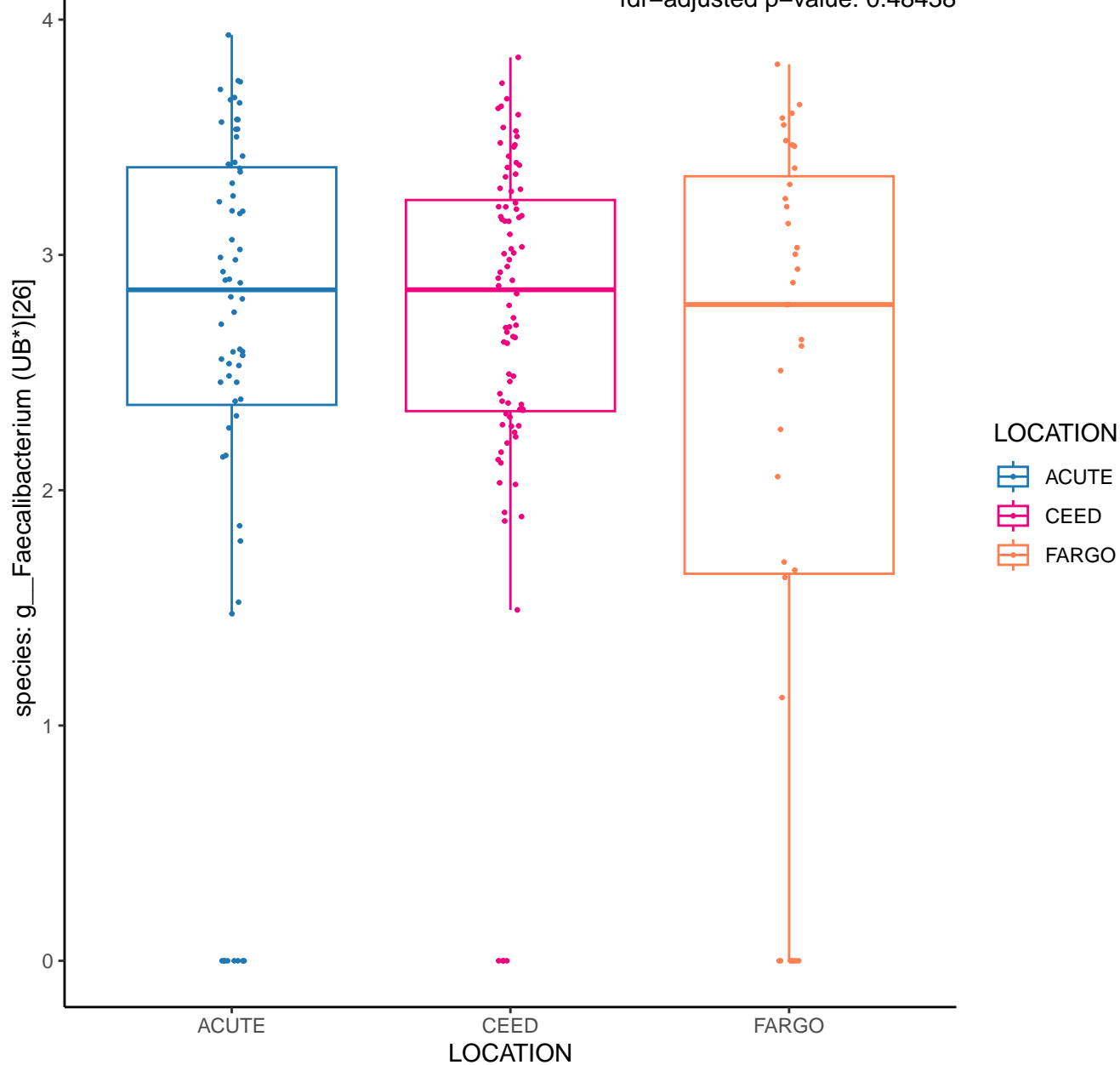
anova p-value: 0.15241

fdr-adjusted p-value: 0.48438



anova p-value: 0.15489

fdr-adjusted p-value: 0.48438



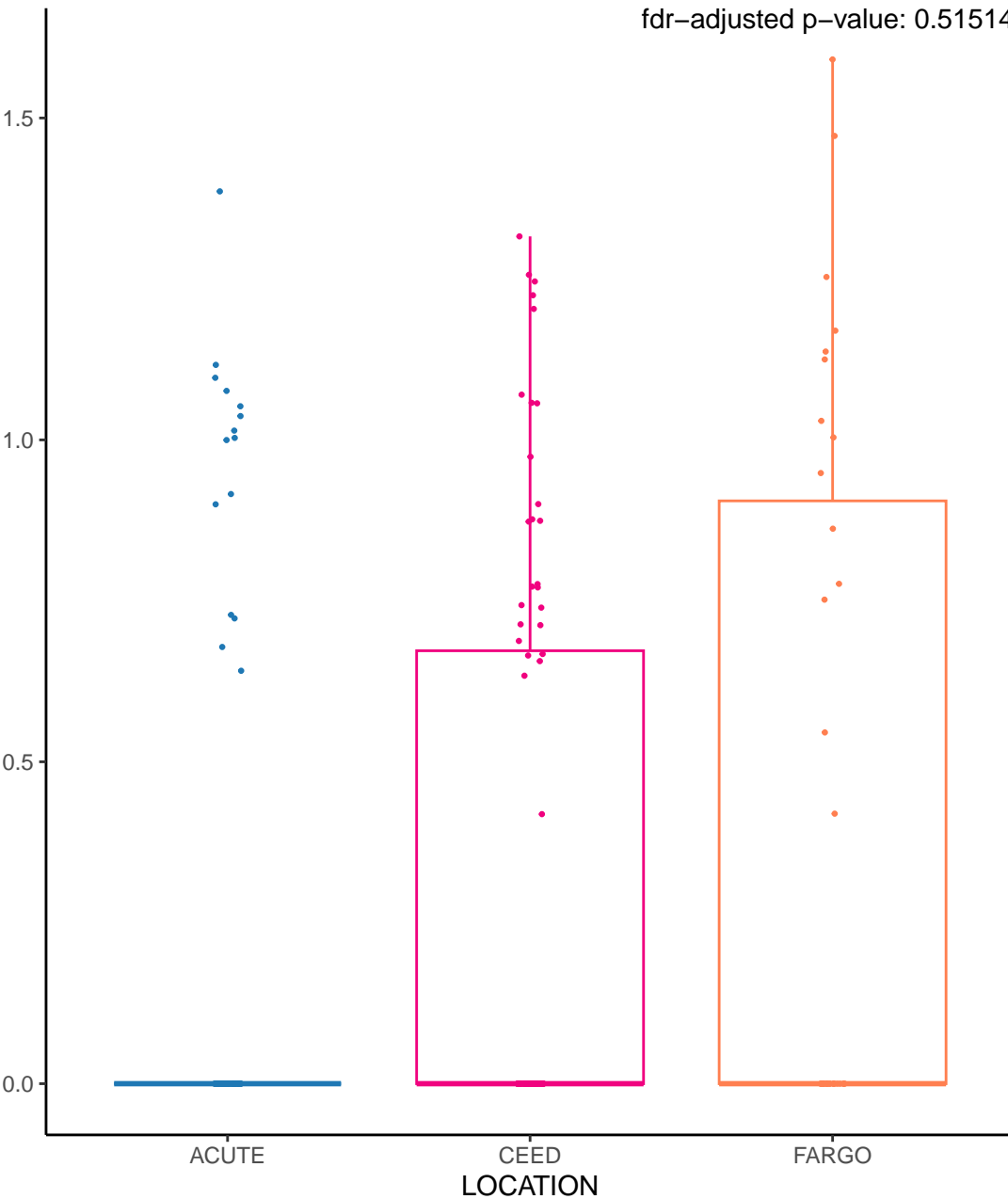
anova p-value: 0.16772

fdr-adjusted p-value: 0.51514

species: Holdemania.[117]

LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.17176

fdr-adjusted p-value: 0.51828

species: uncultured_organism[59]

LOCATION

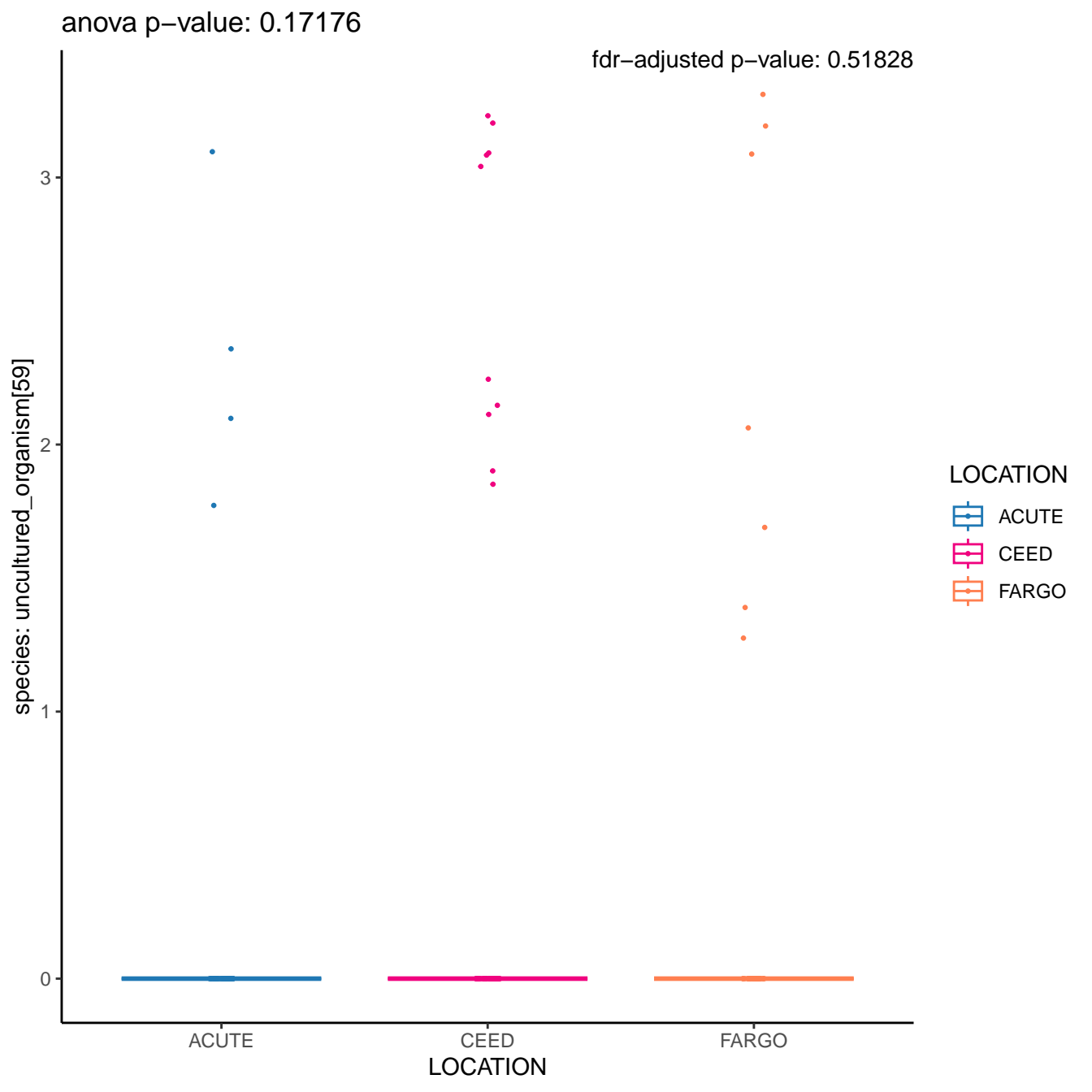
ACUTE
CEED
FARGO

ACUTE

CEED

FARGO

LOCATION



anova p-value: 0.18341

fdr-adjusted p-value: 0.54391

species: *Desulfovibrio*. [95]

LOCATION

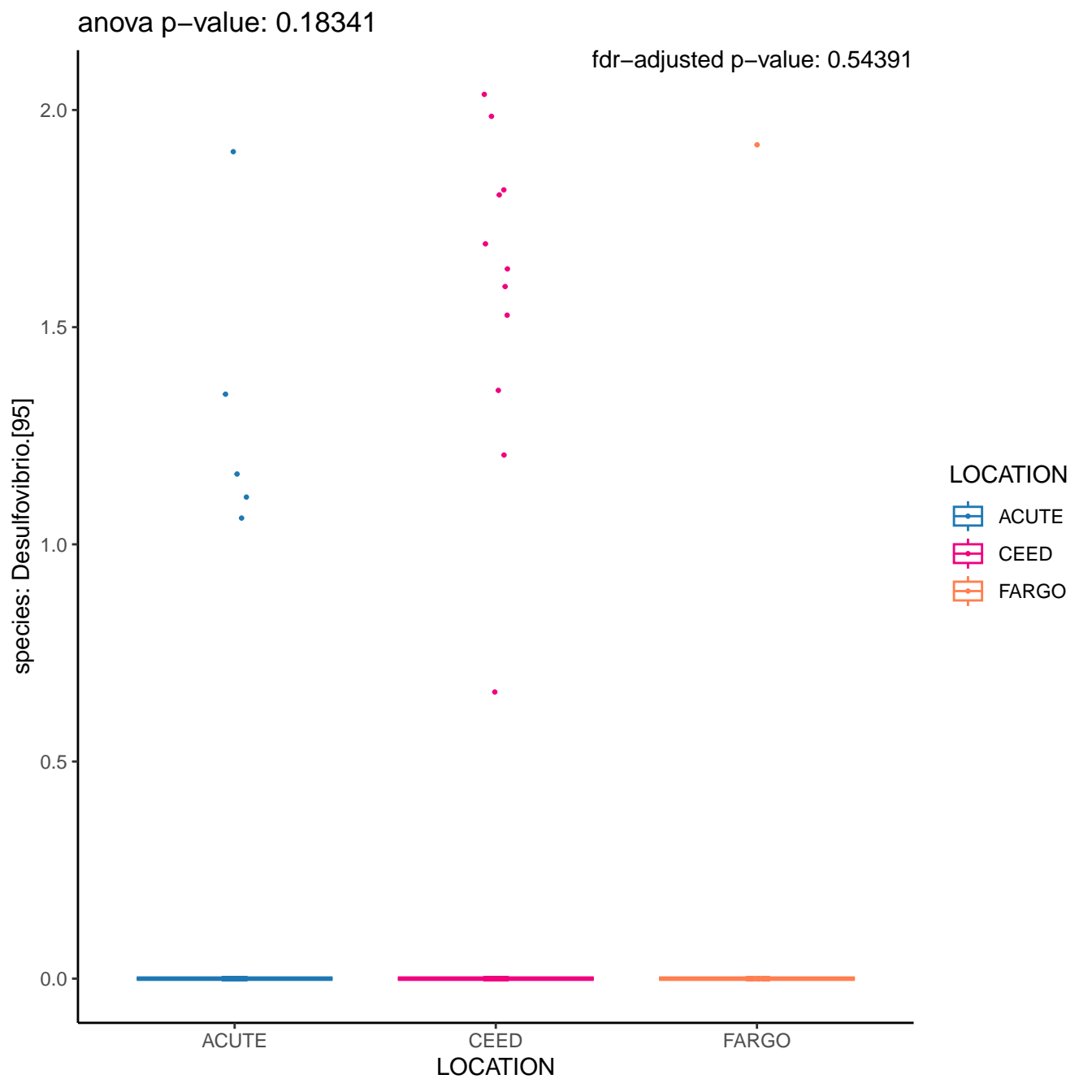
ACUTE
CEED
FARGO

ACUTE

CEED

FARGO

LOCATION

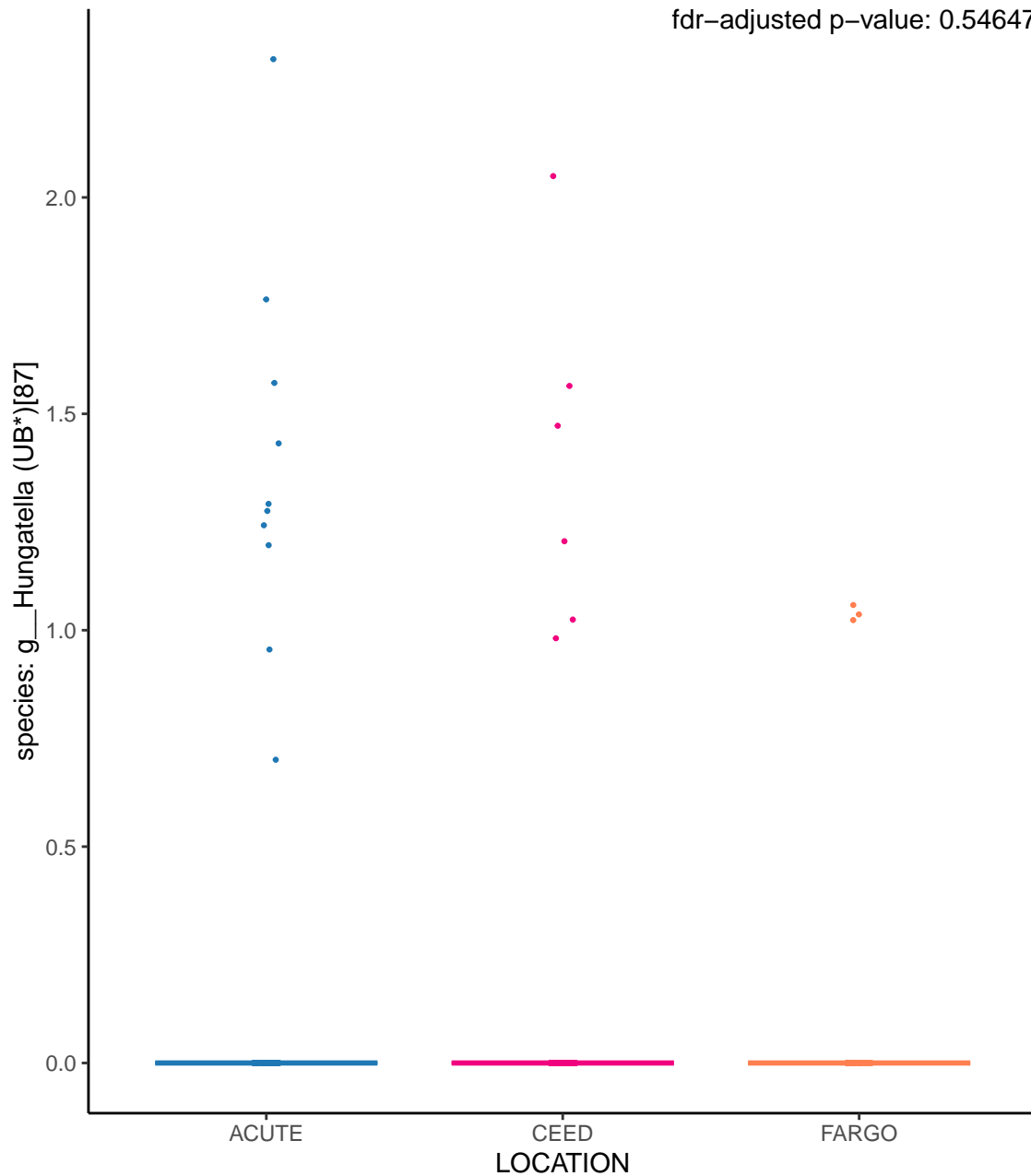
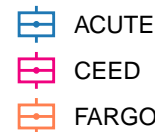


anova p-value: 0.19433

fdr-adjusted p-value: 0.54647

species: g__Hungatella (UB*)[87]

LOCATION



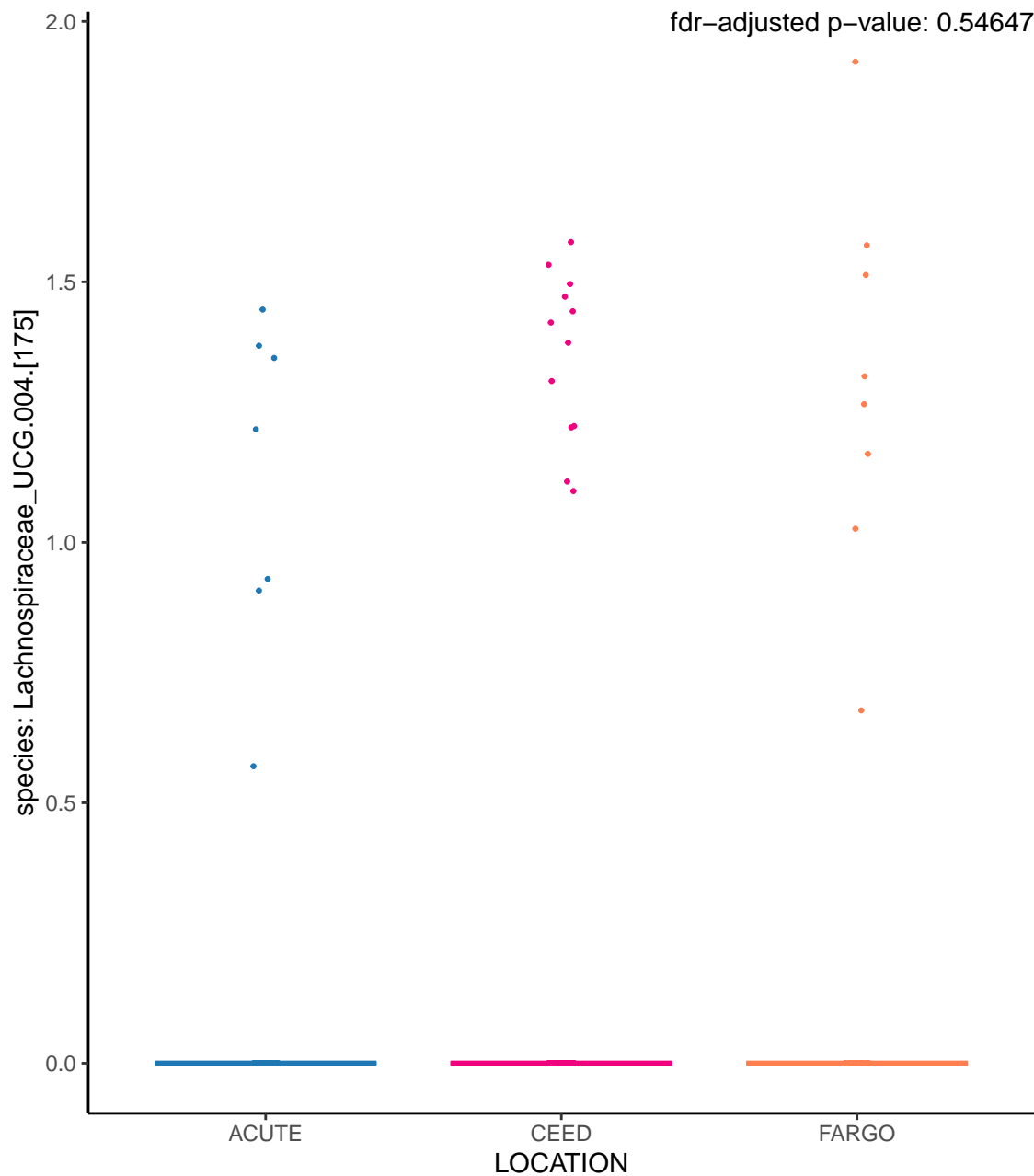
anova p-value: 0.19703

fdr-adjusted p-value: 0.54647

species: Lachnospiraceae_UCG.004.[175]

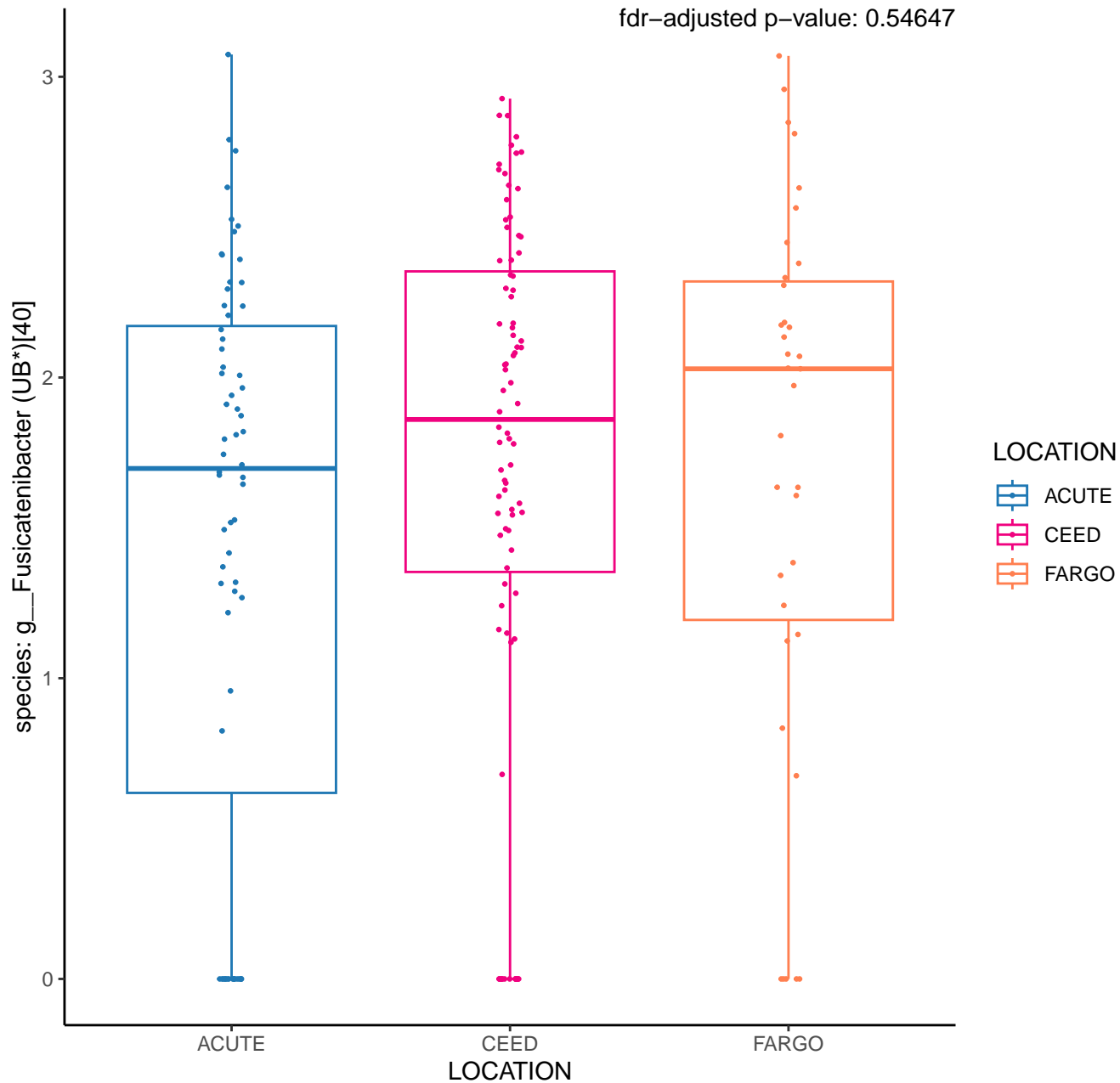
LOCATION

ACUTE
CEED
FARGO



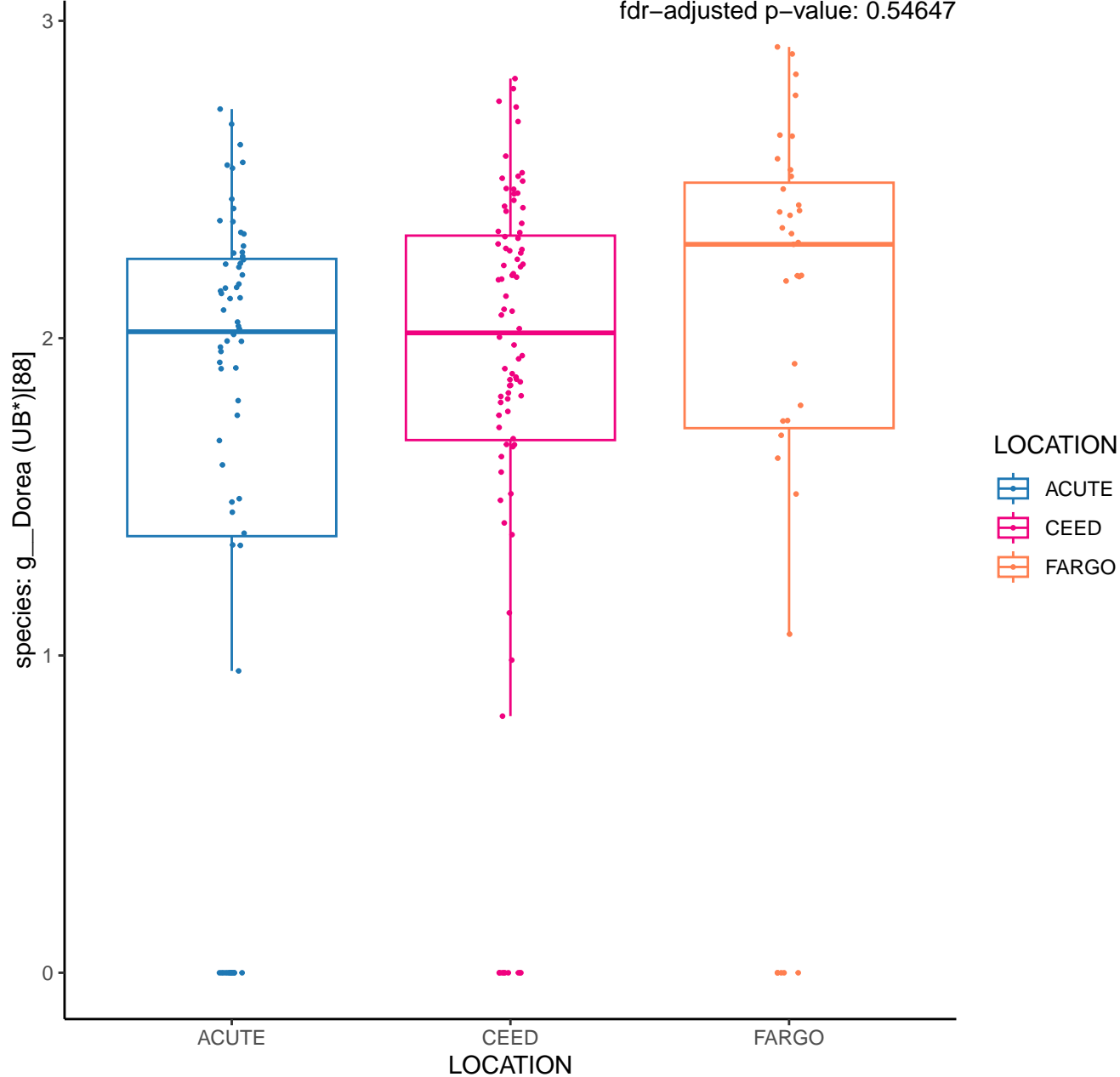
anova p-value: 0.19842

fdr-adjusted p-value: 0.54647



anova p-value: 0.19994

fdr-adjusted p-value: 0.54647



anova p-value: 0.20398

fdr-adjusted p-value: 0.54647

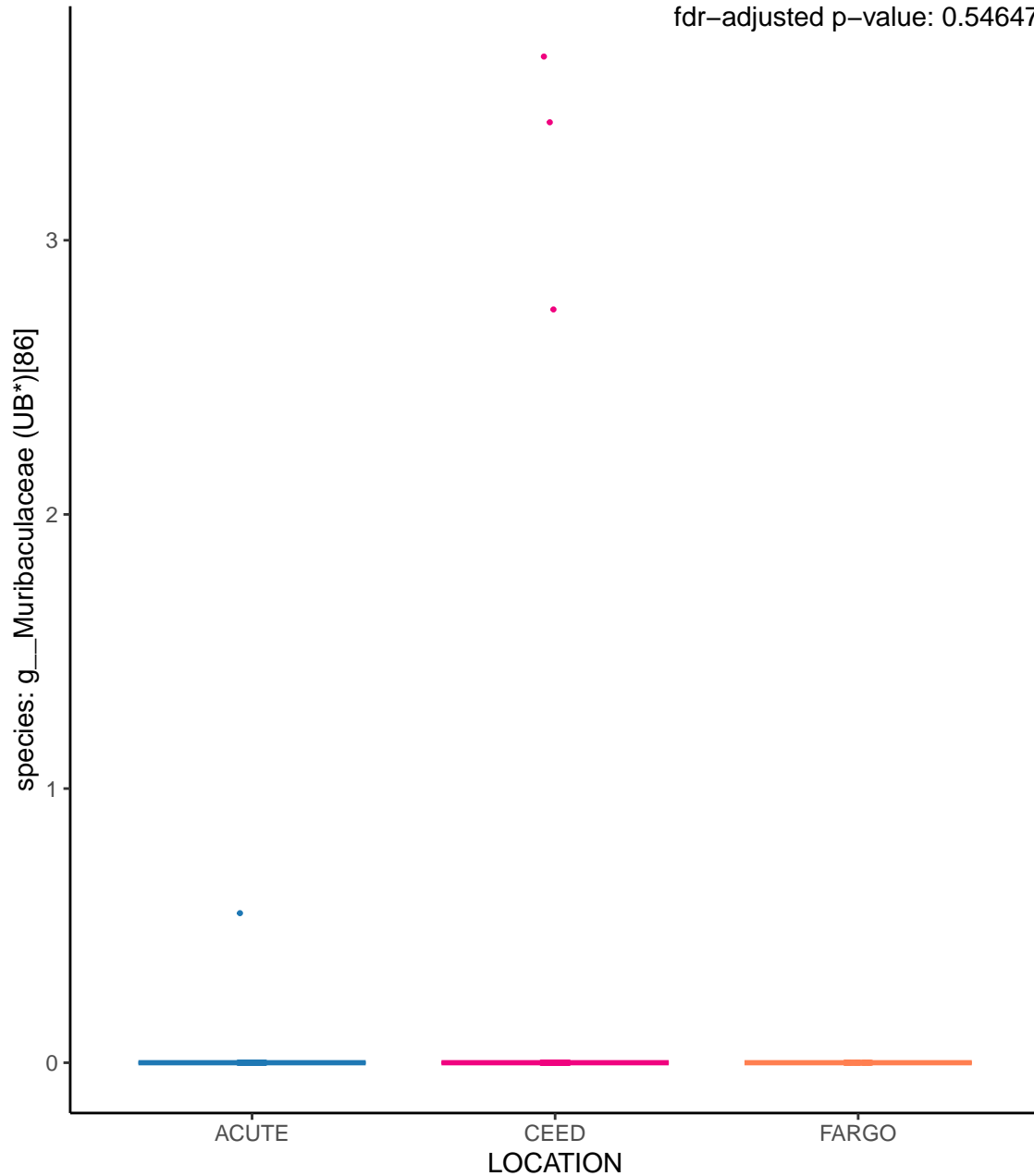
species: g__Muribaculaceae (UB*)[86]

LOCATION

ACUTE

CEED

FARGO



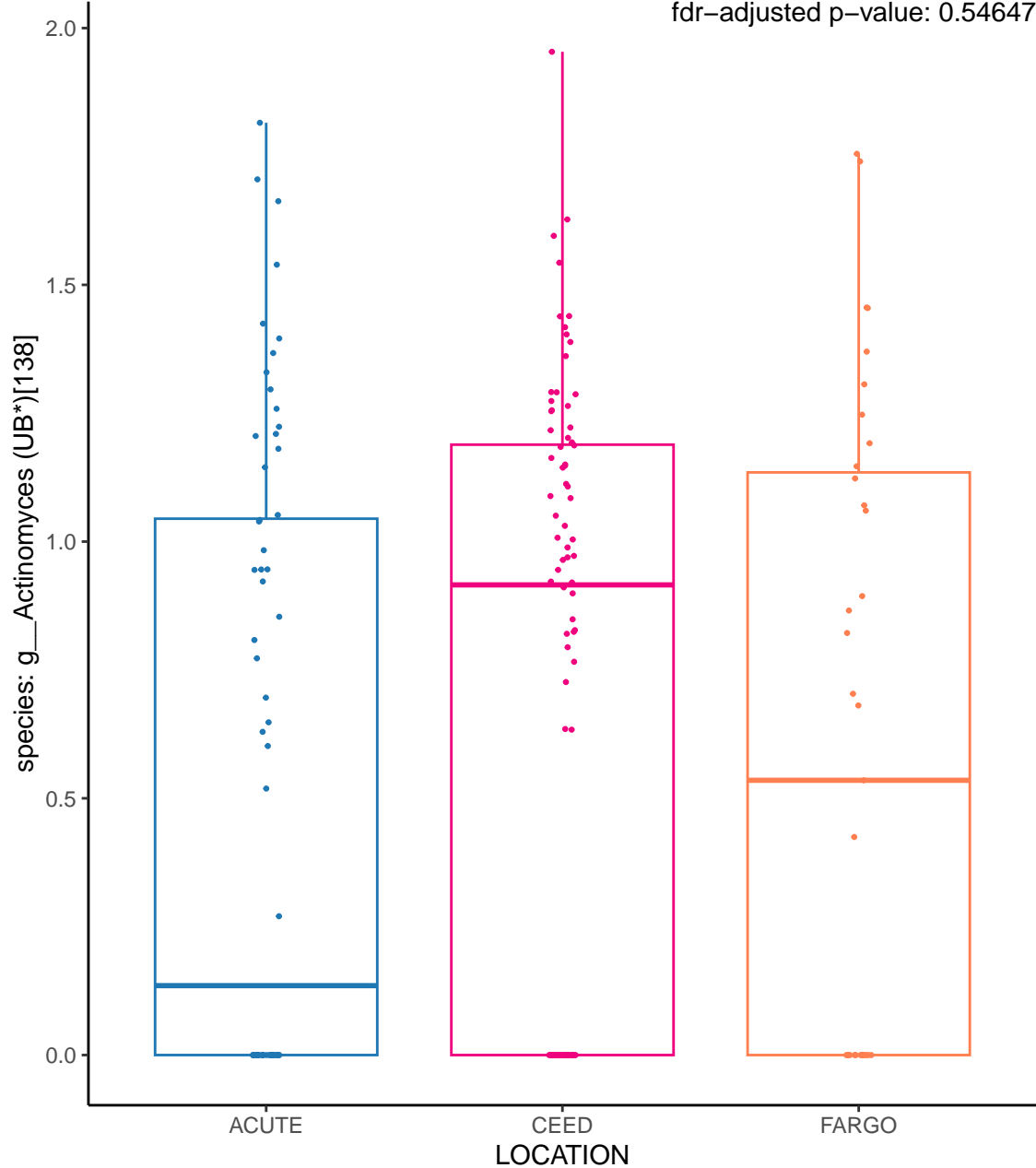
anova p-value: 0.20484

fdr-adjusted p-value: 0.54647

species: g__Actinomyces (UB*)[138]

LOCATION

ACUTE
CEED
FARGO



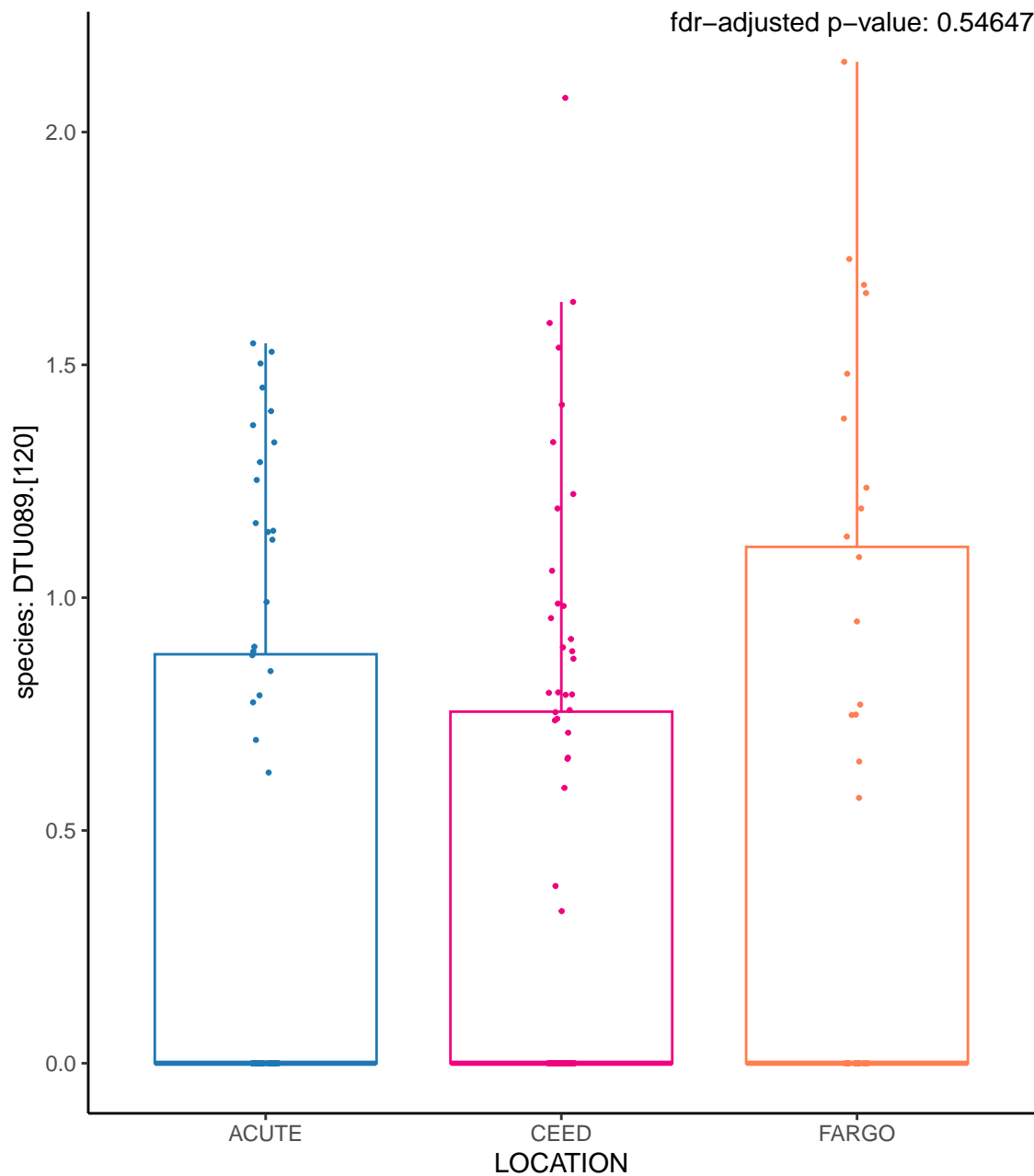
anova p-value: 0.20651

fdr-adjusted p-value: 0.54647

species: DTU089.[120]

LOCATION

ACUTE
CEED
FARGO

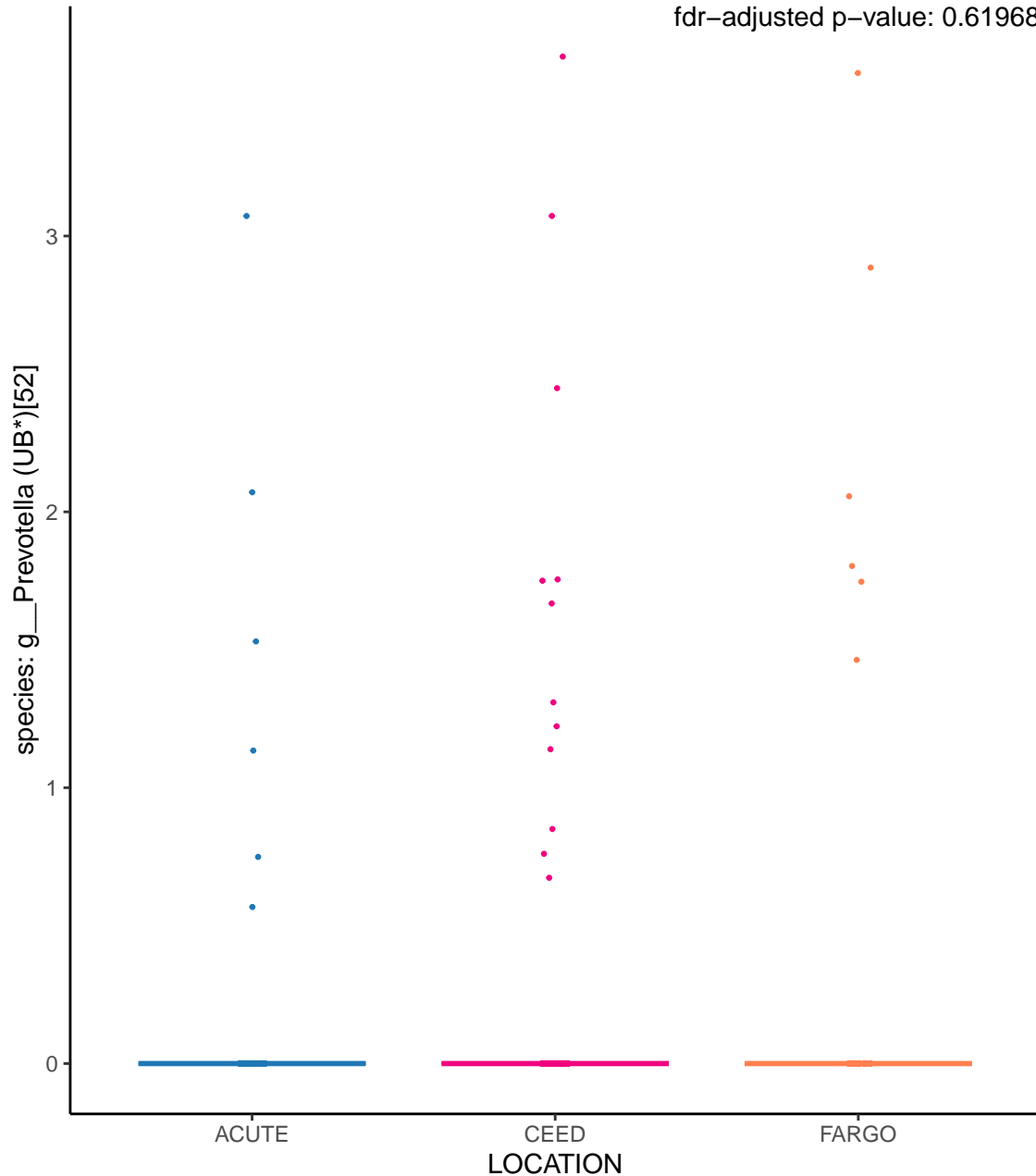


anova p-value: 0.23832

fdr-adjusted p-value: 0.61968

species: g__Prevotella (UB*)[52]

LOCATION



anova p-value: 0.24264

fdr-adjusted p-value: 0.61968

species: Bifidobacterium_dentium[51]

LOCATION

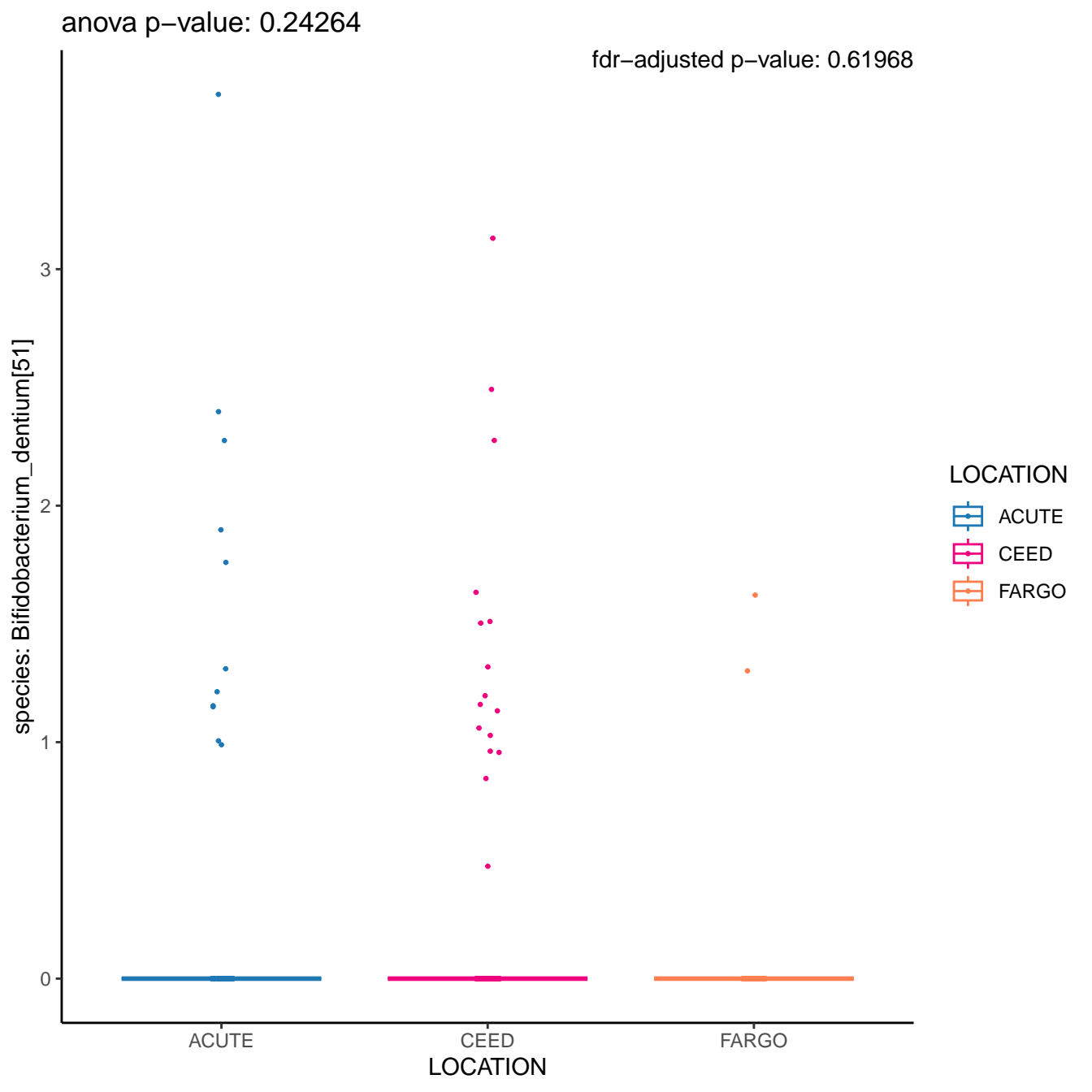


ACUTE

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FARGO

LOCATION



anova p-value: 0.24499

fdr-adjusted p-value: 0.61968

species: Enterococcus.[150]

LOCATION

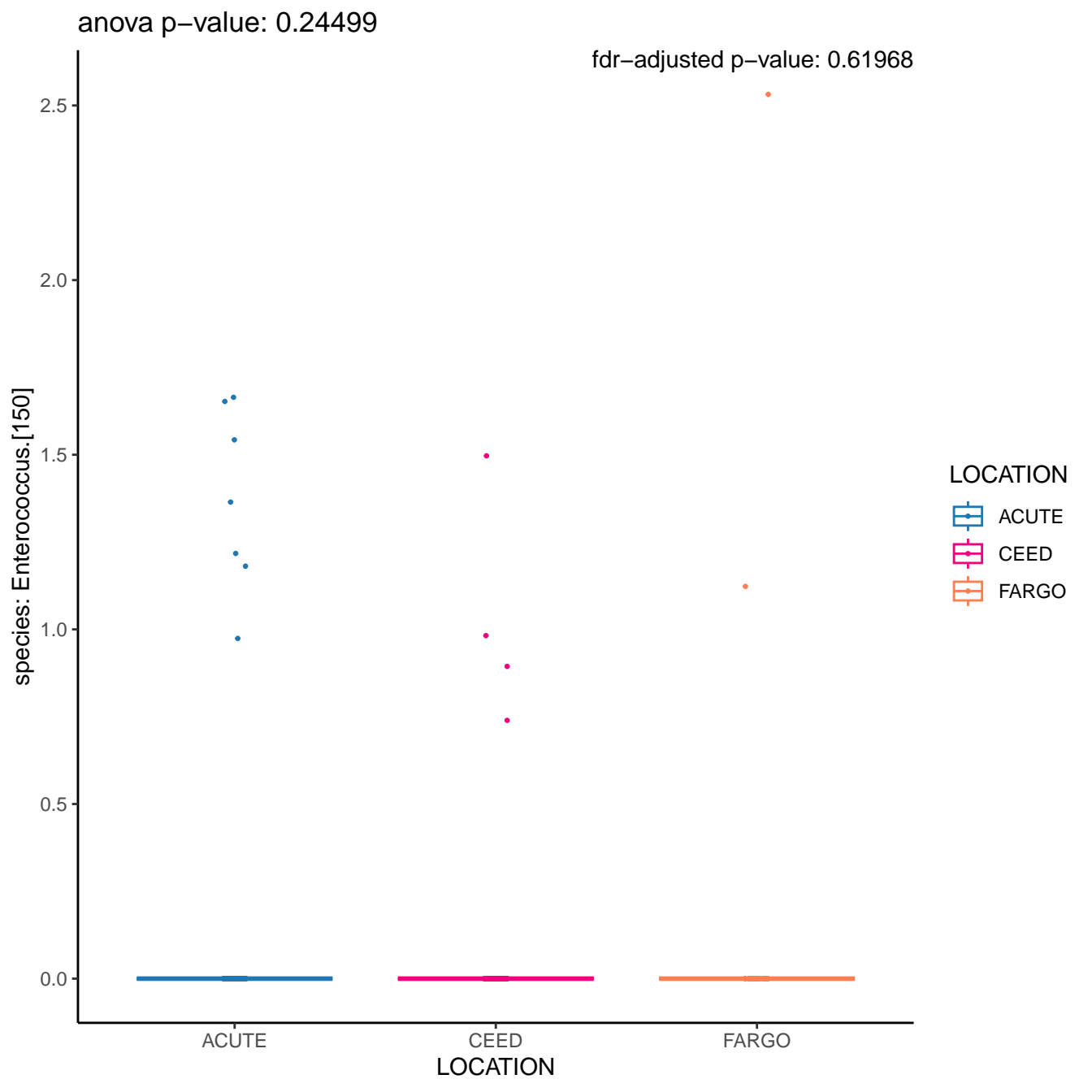
ACUTE
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ACUTE

CEED

FARGO

LOCATION



anova p-value: 0.24892

fdr-adjusted p-value: 0.62048

species: uncultured.[98]

LOCATION

ACUTE

CEED

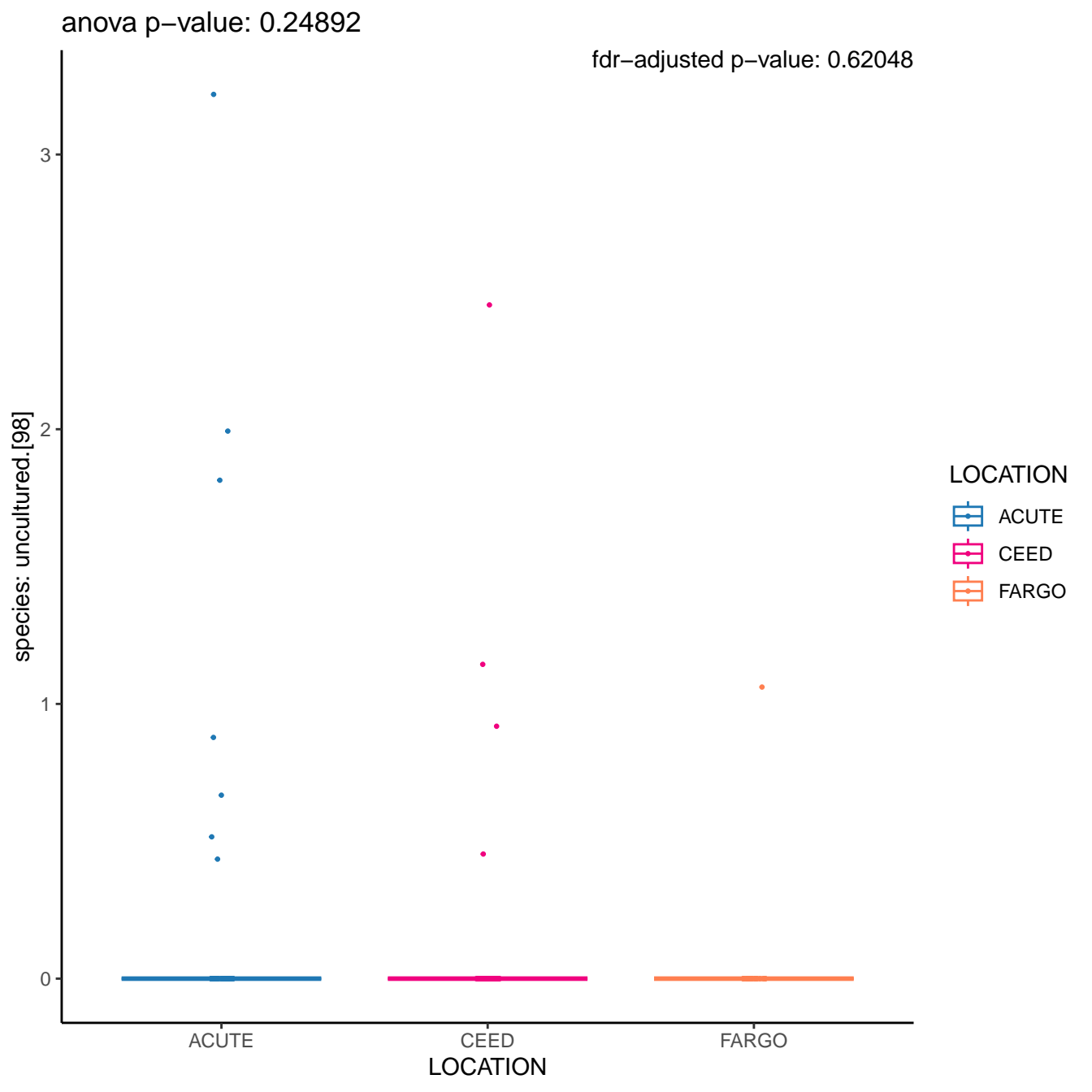
FARGO

ACUTE

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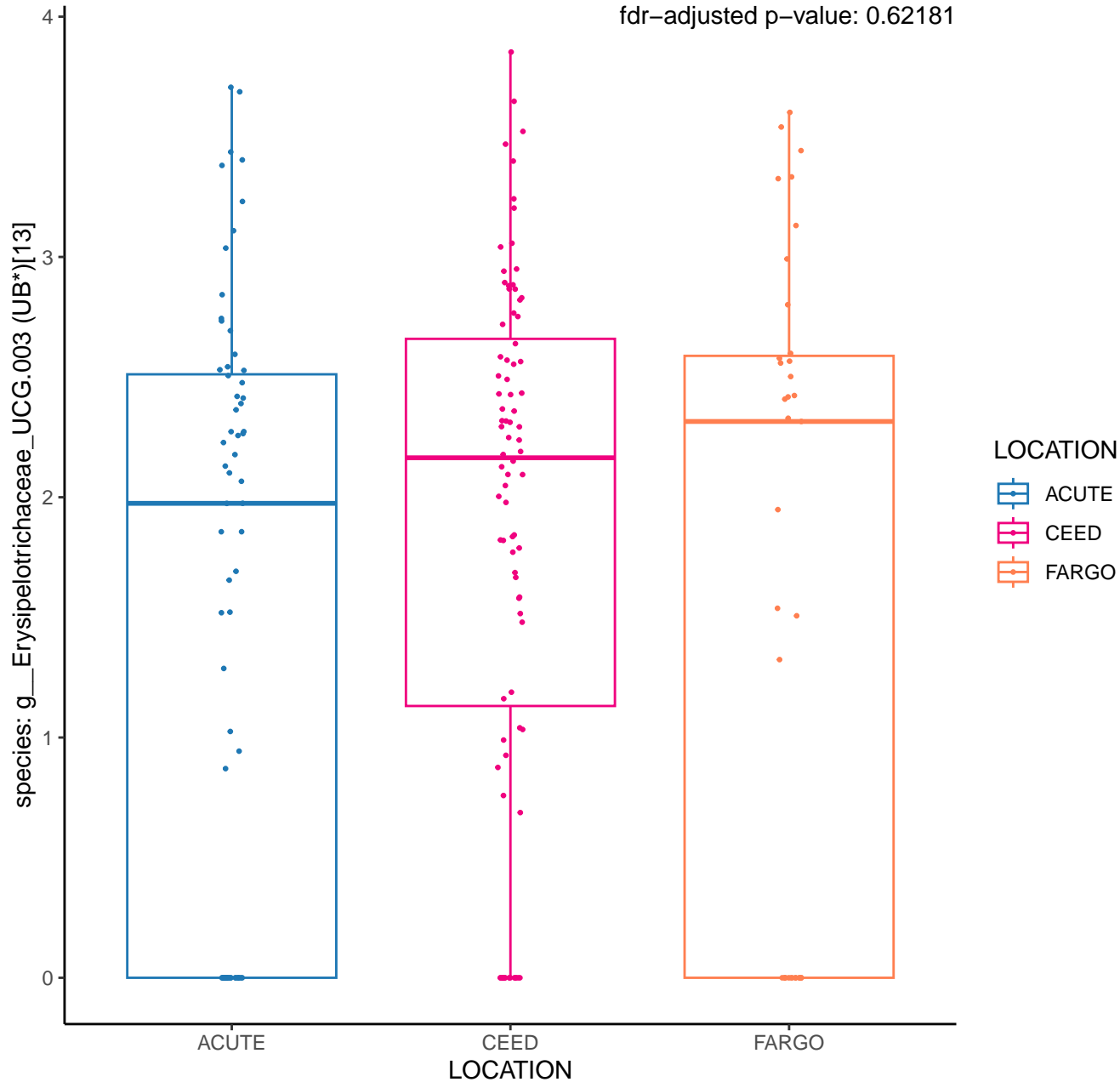
FARGO

LOCATION



anova p-value: 0.25475

fdr-adjusted p-value: 0.62181



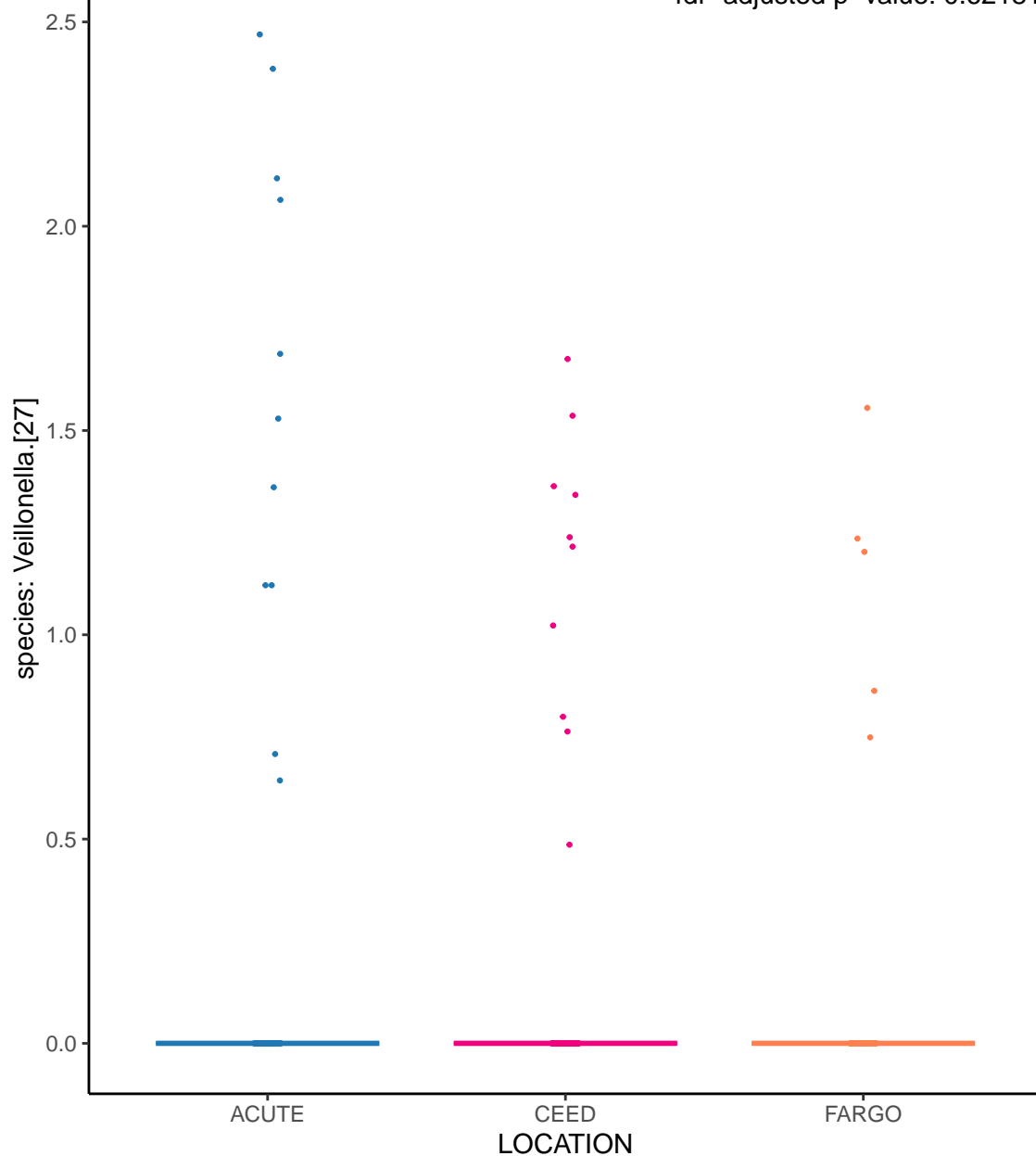
anova p-value: 0.26373

fdr-adjusted p-value: 0.62181

species: *Veillonella*. [27]

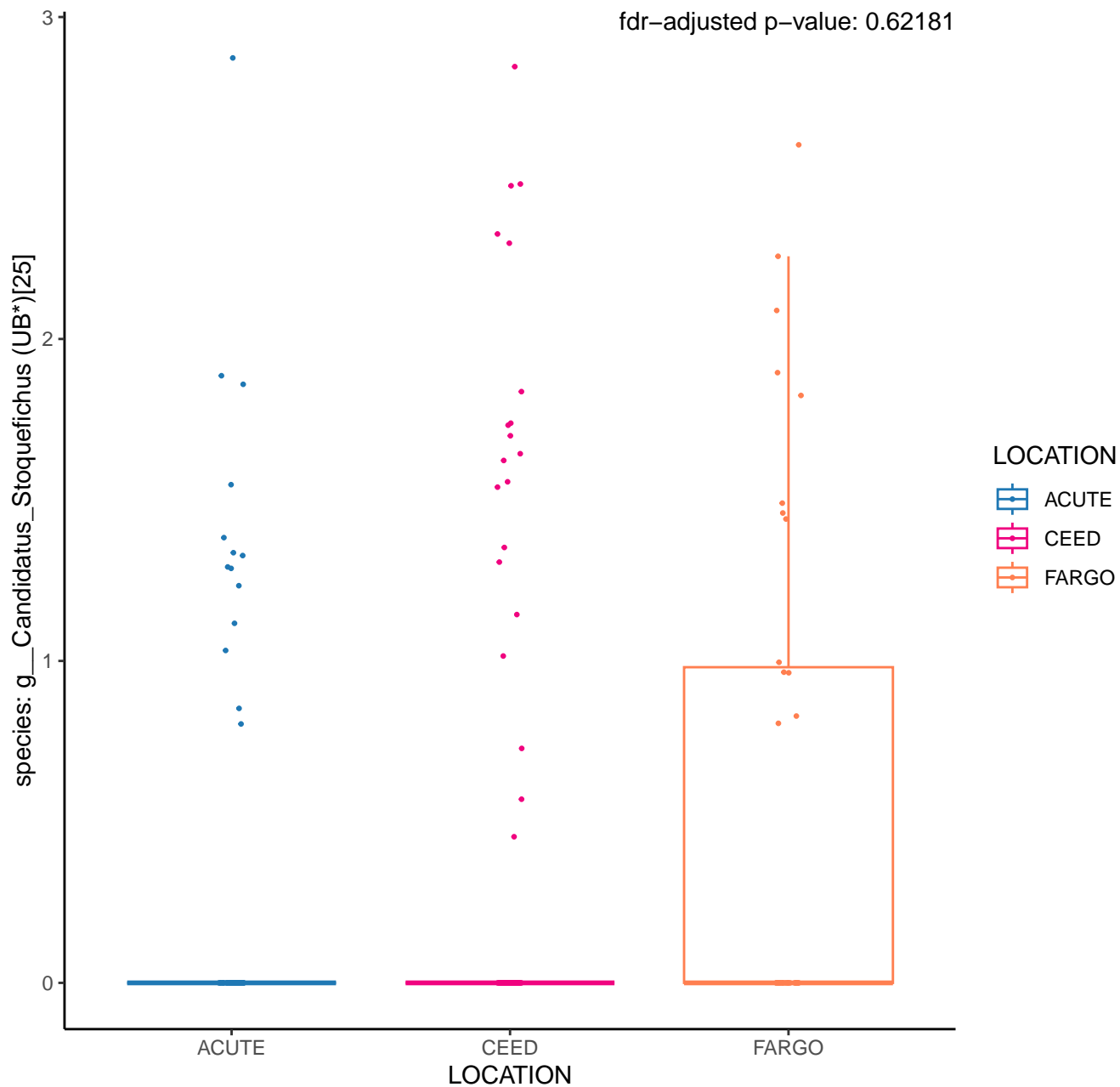
LOCATION

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FARGO

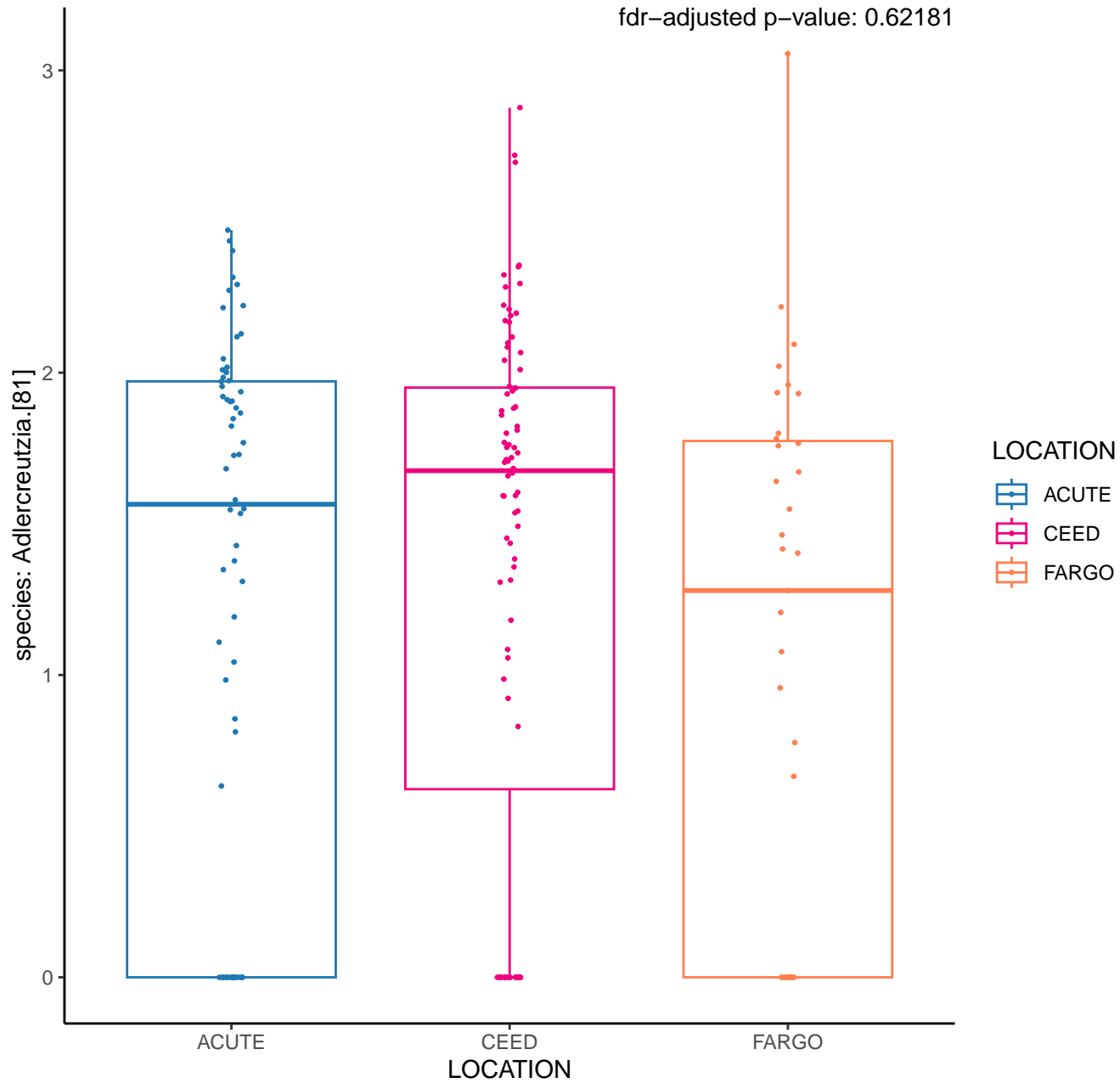


anova p-value: 0.26529

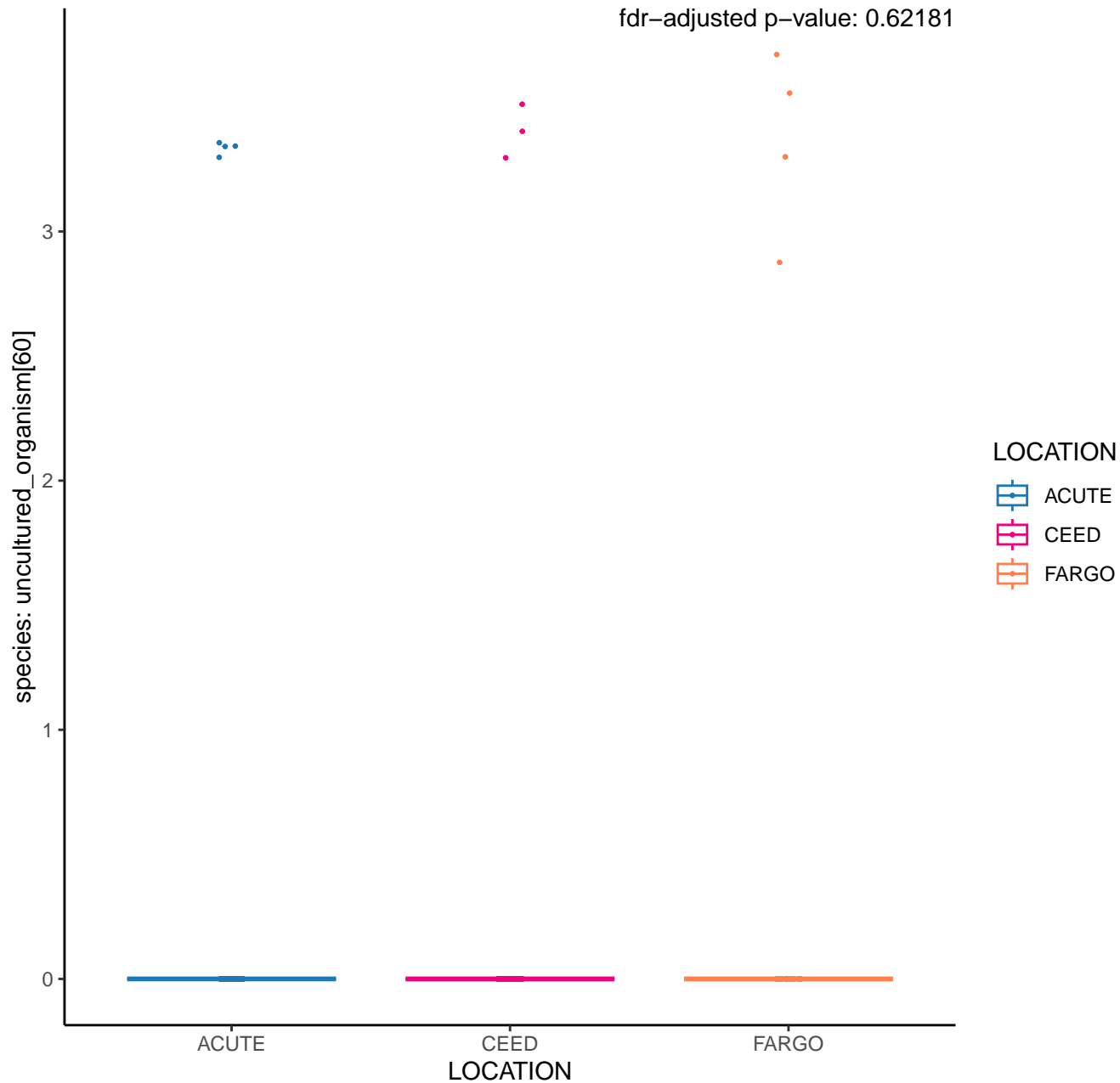
fdr-adjusted p-value: 0.62181



fdr-adjusted p-value: 0.62181



fdr-adjusted p-value: 0.62181



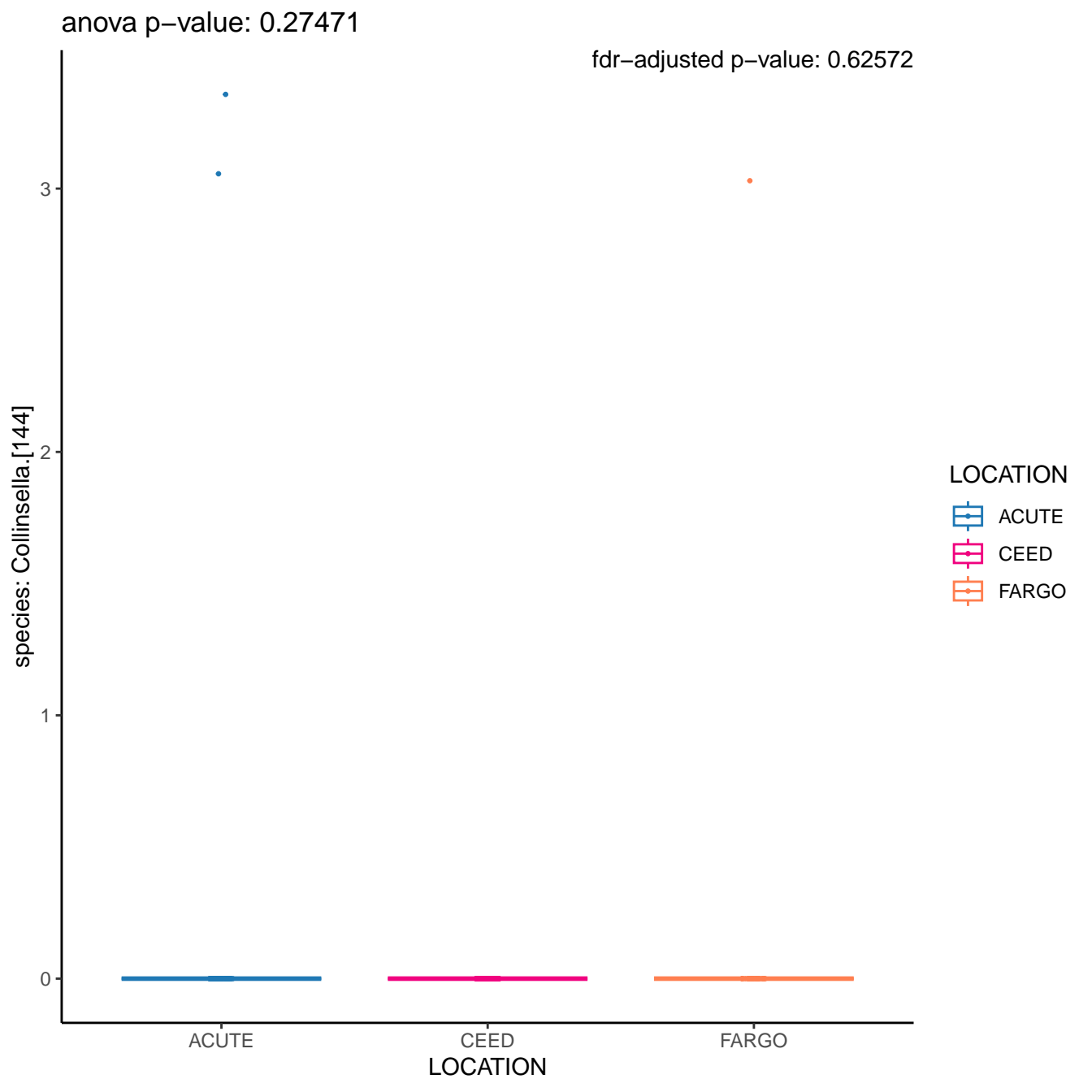
anova p-value: 0.27471

fdr-adjusted p-value: 0.62572

species: Collinsella.[144]

LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.27851

fdr-adjusted p-value: 0.62572

species: *Lactobacillus_helveticus*[142]

LOCATION

ACUTE

CEED

FARGO

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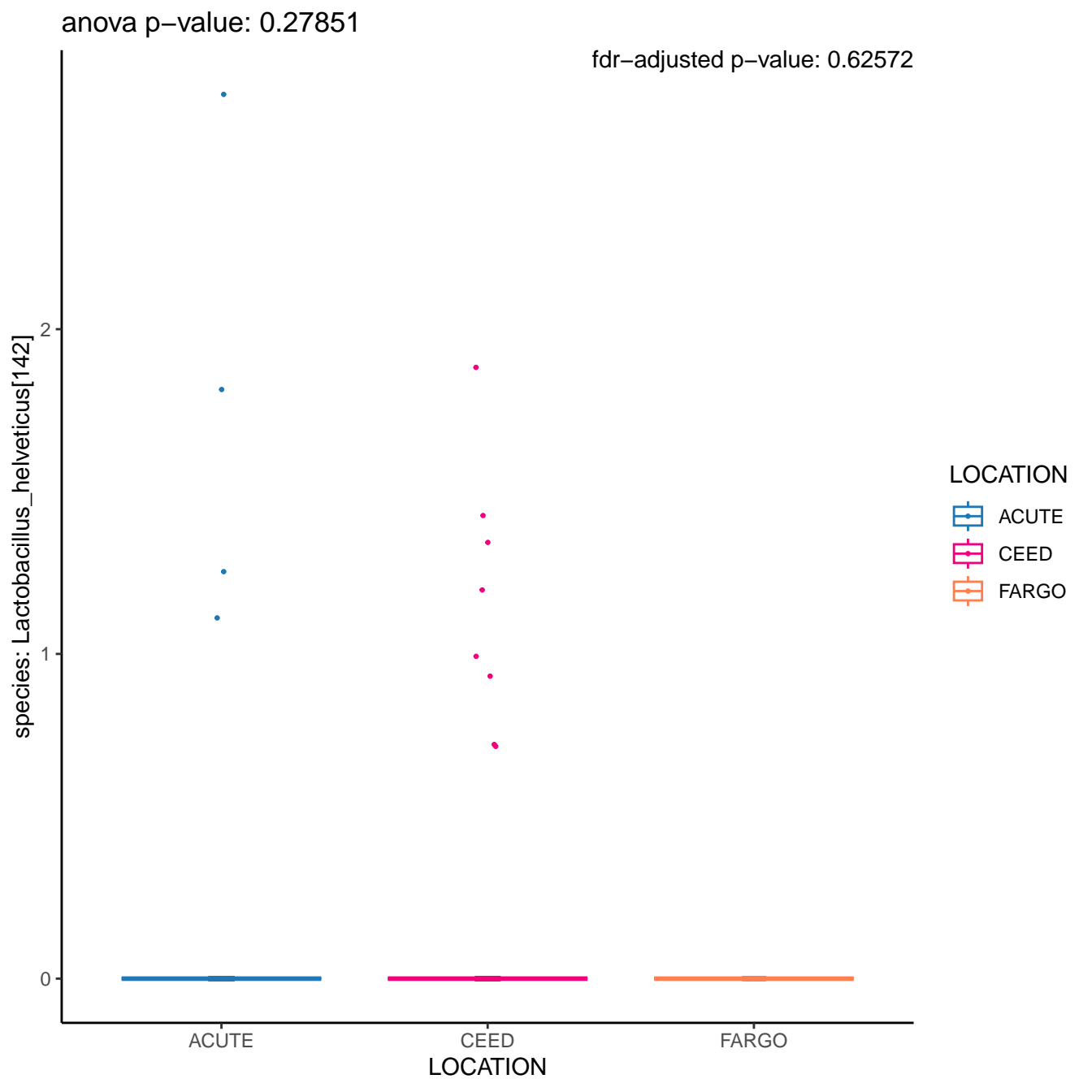
2

ACUTE

CEED

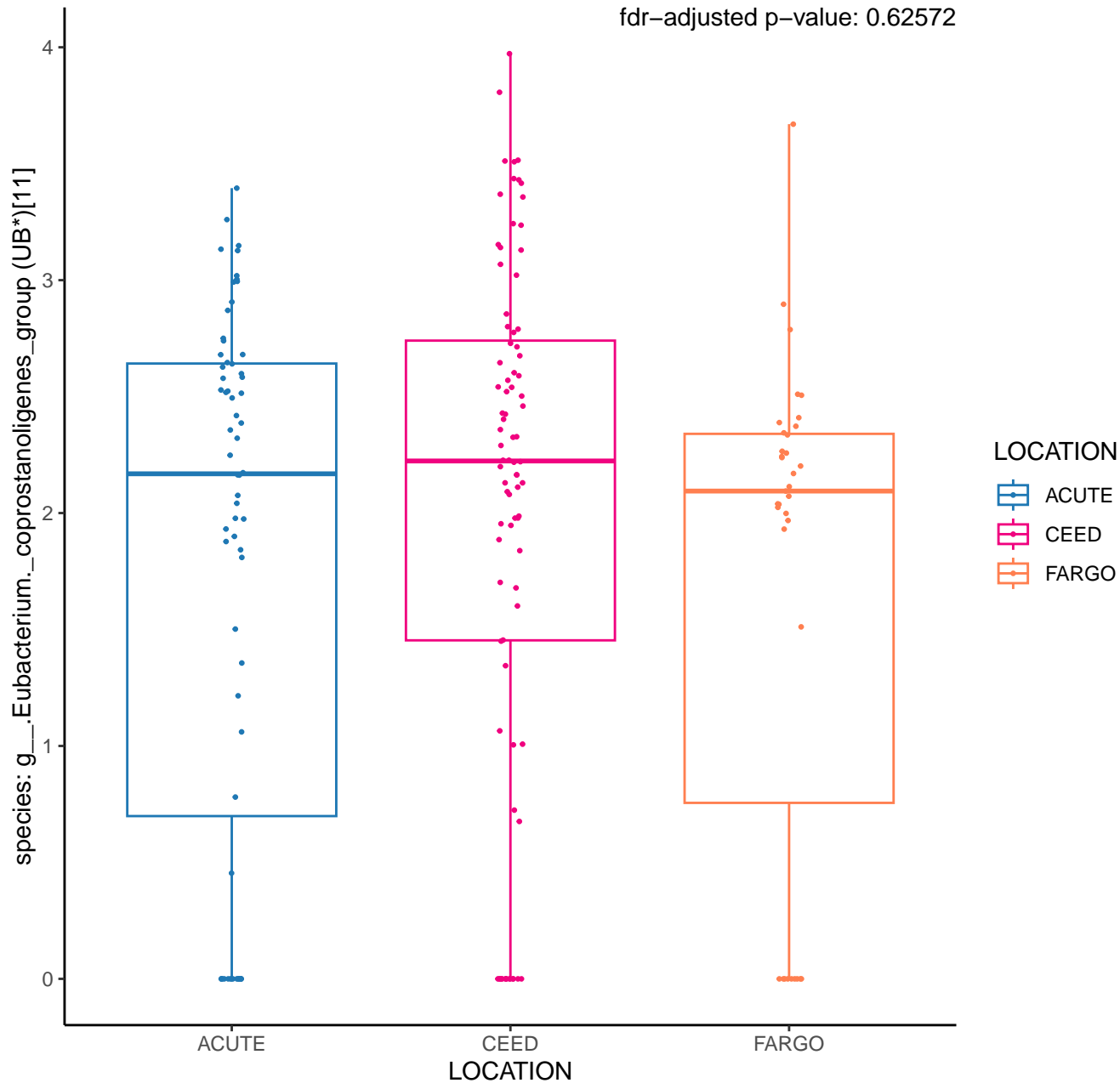
FARGO

LOCATION



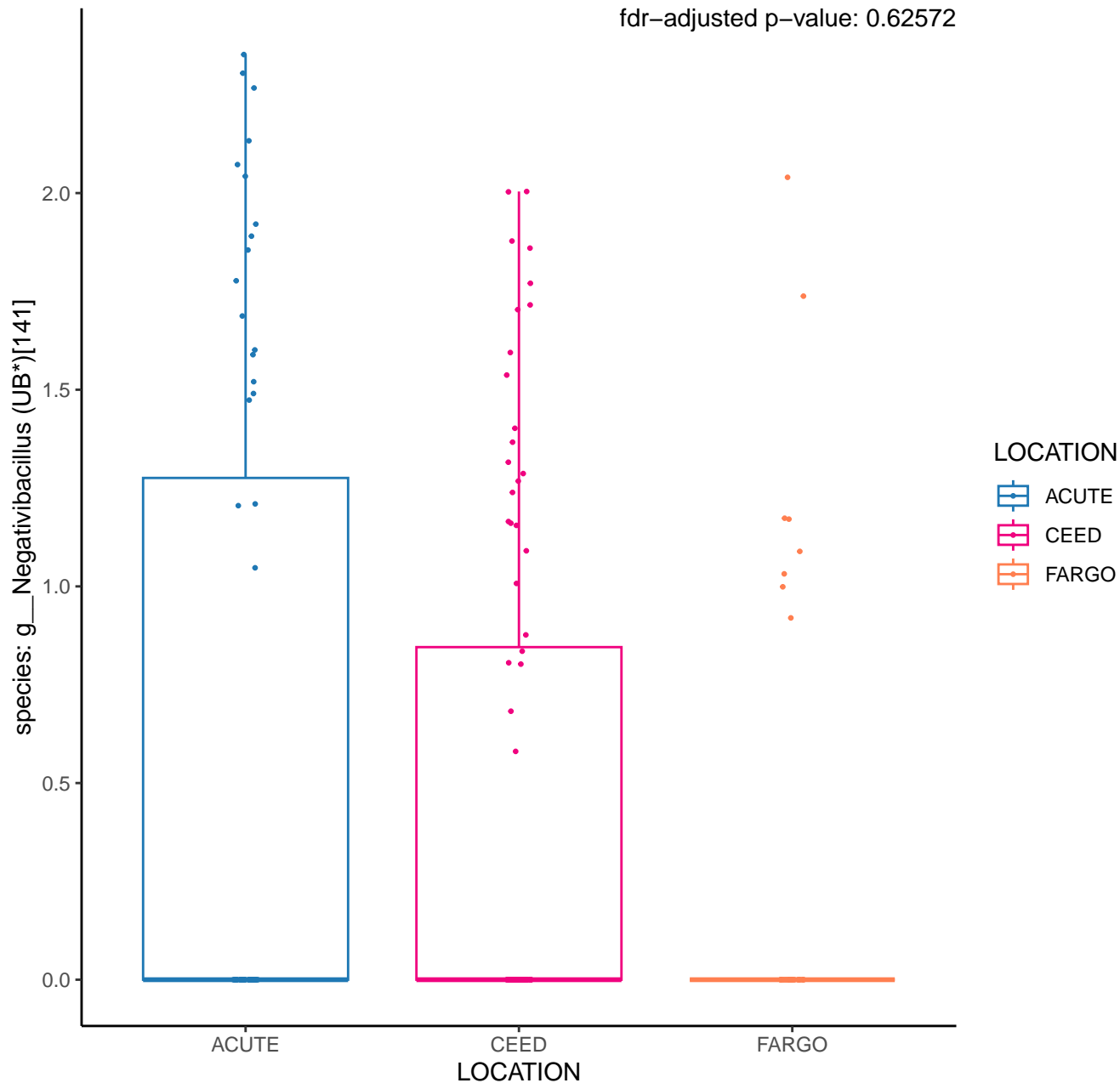
anova p-value: 0.28334

fdr-adjusted p-value: 0.62572



anova p-value: 0.28376

fdr-adjusted p-value: 0.62572



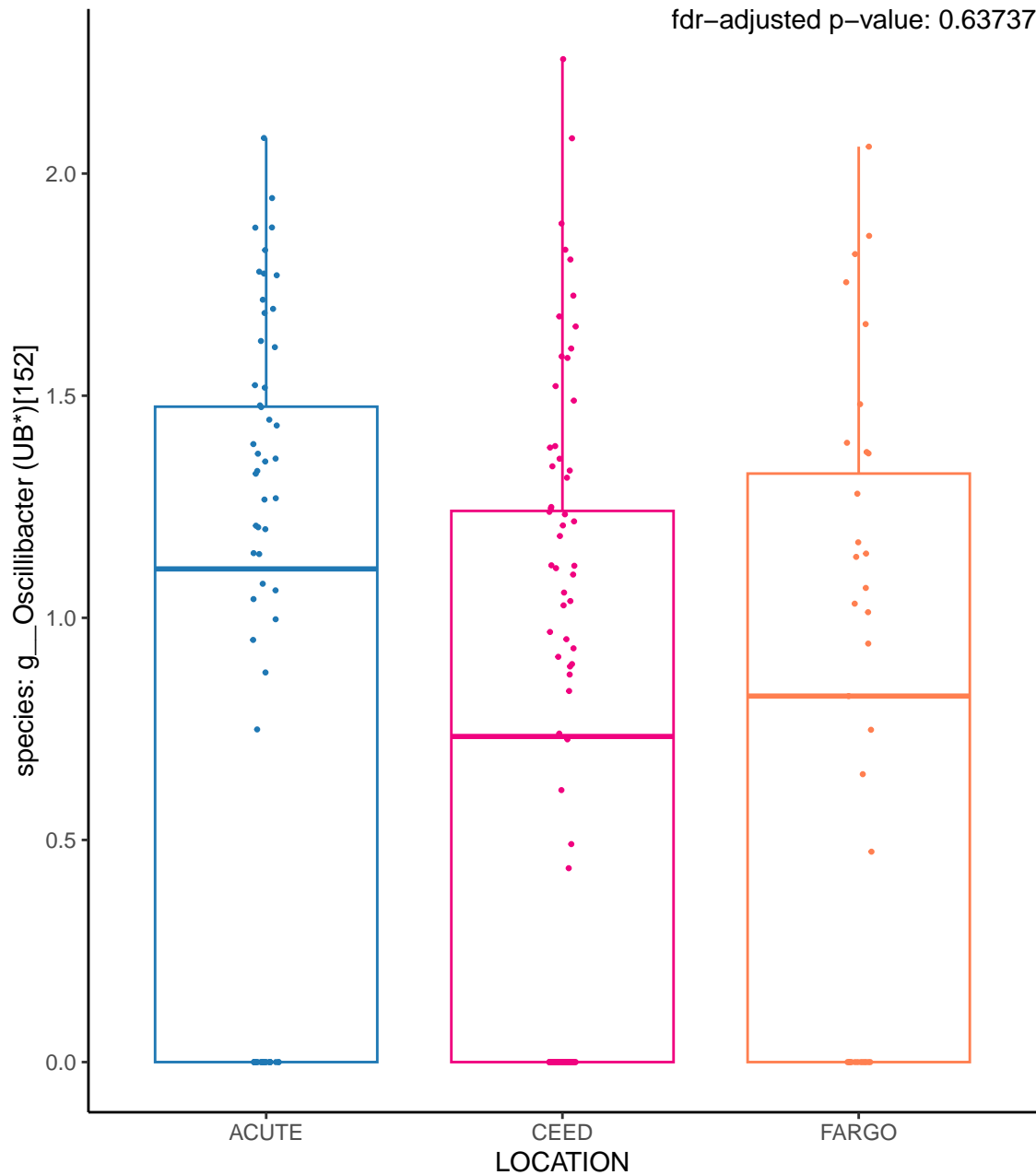
anova p-value: 0.29274

fdr-adjusted p-value: 0.63737

species: g__Oscillibacter (UB*)[152]

LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.30366

fdr-adjusted p-value: 0.65288

species: Ruminococcus.[99]

LOCATION

ACUTE

CEED

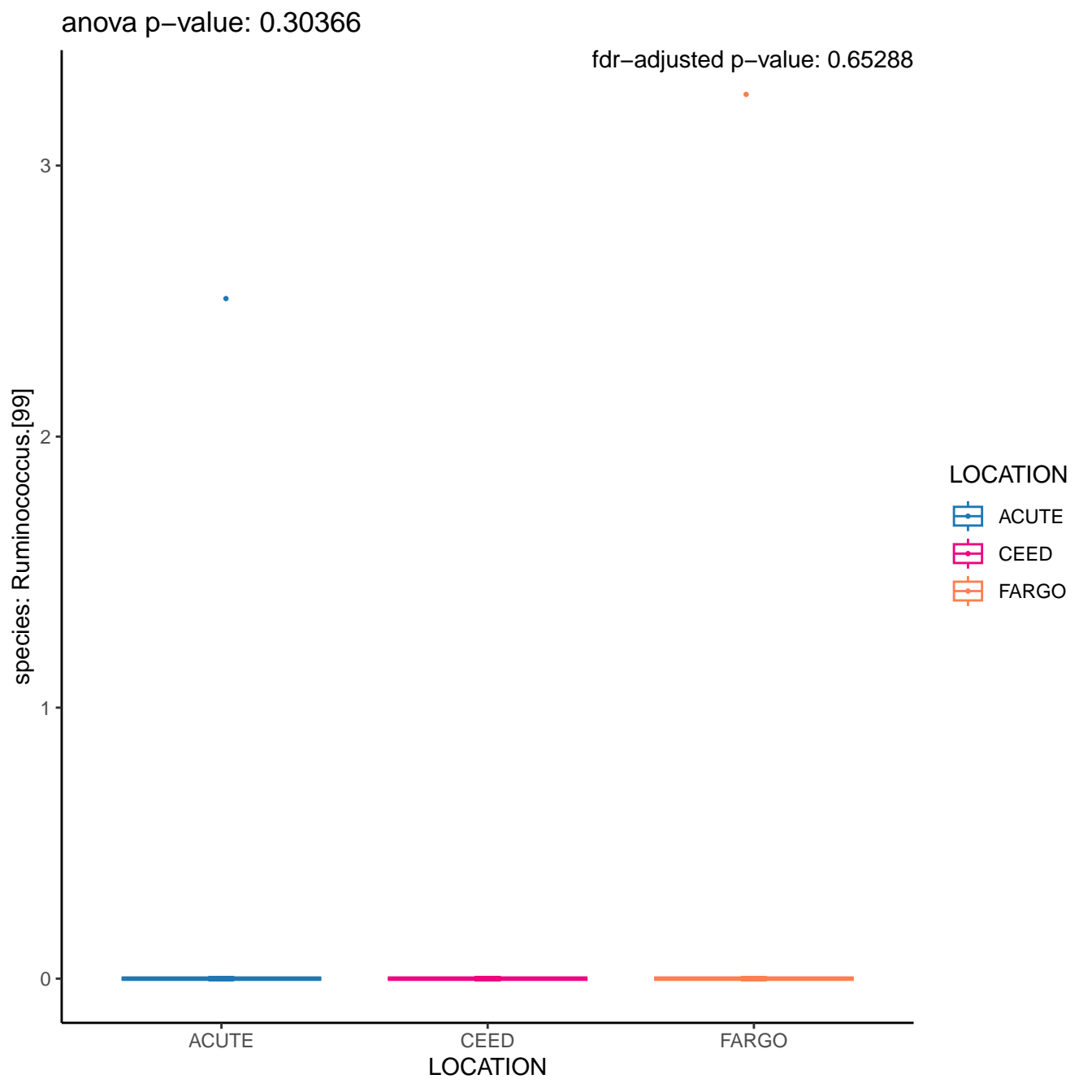
FARGO

LOCATION

ACUTE

CEED

FARGO



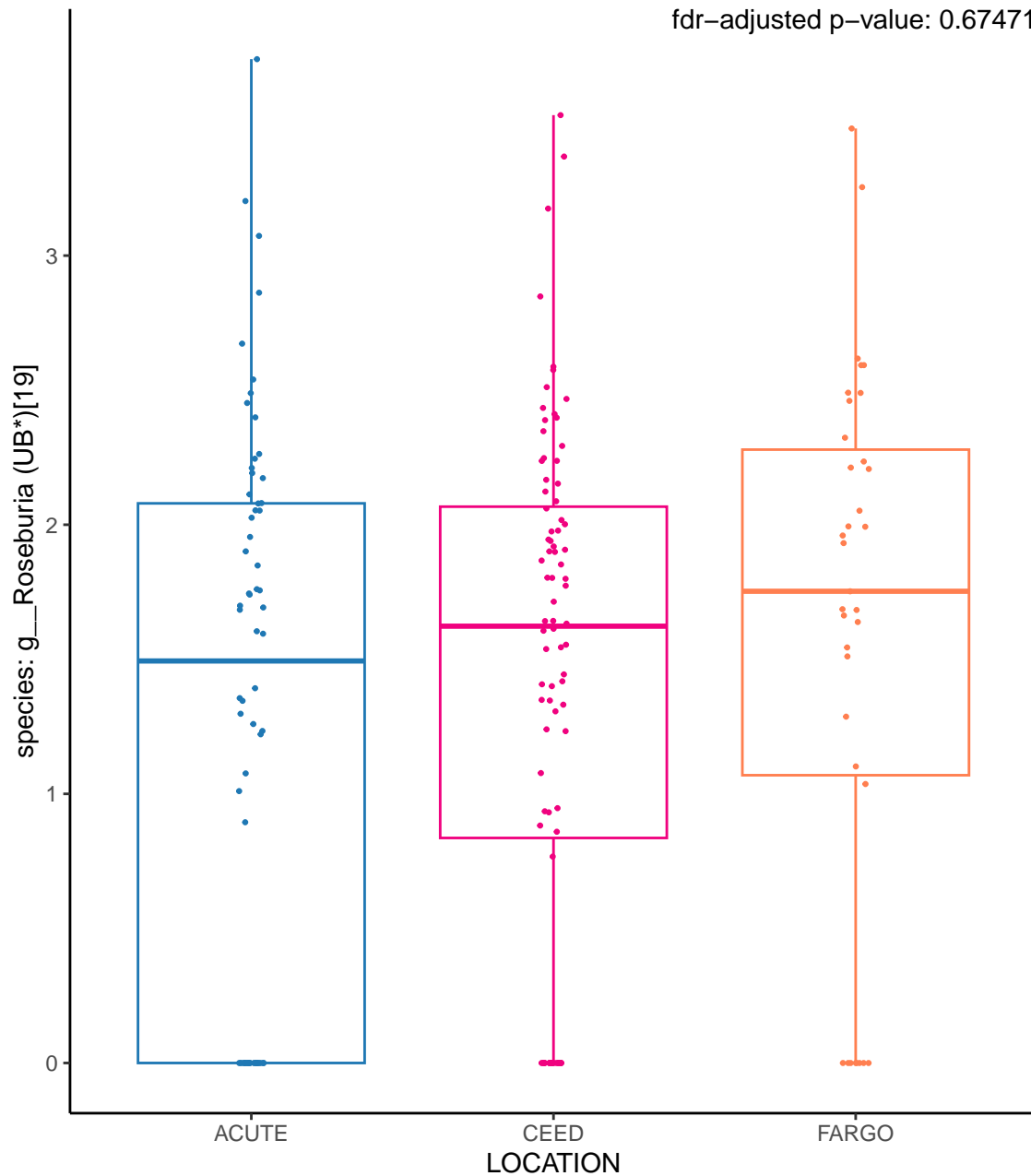
anova p-value: 0.32637

fdr-adjusted p-value: 0.67471

species: g__Roseburia (UB*)[19]

LOCATION

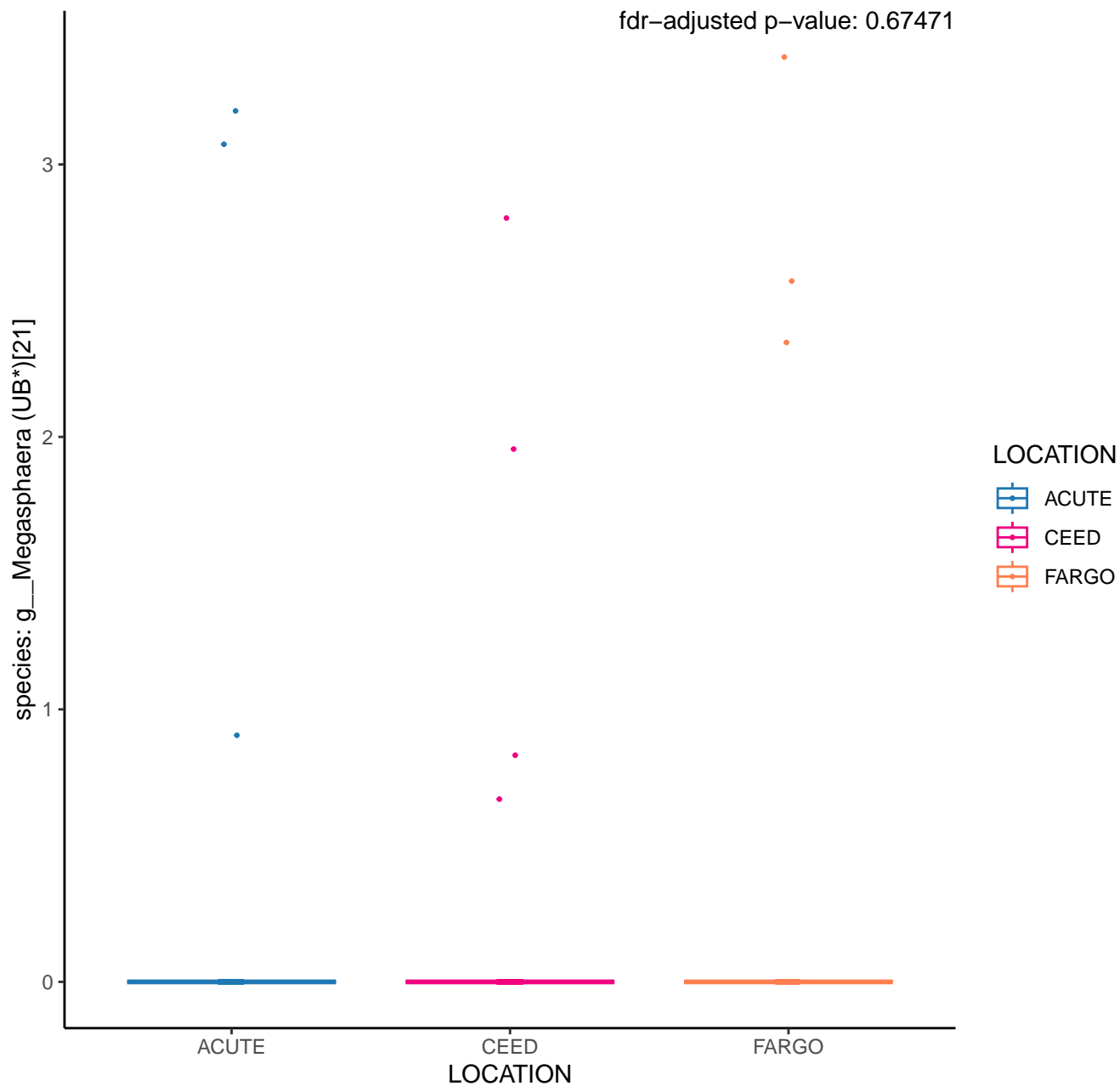
ACUTE
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anova p-value: 0.33277

fdr-adjusted p-value: 0.67471

species: g__Megasphaera (UB*)[21]



anova p-value: 0.33506

fdr-adjusted p-value: 0.67471

species: *Escherichia_coli*[16]

LOCATION

ACUTE

CEED

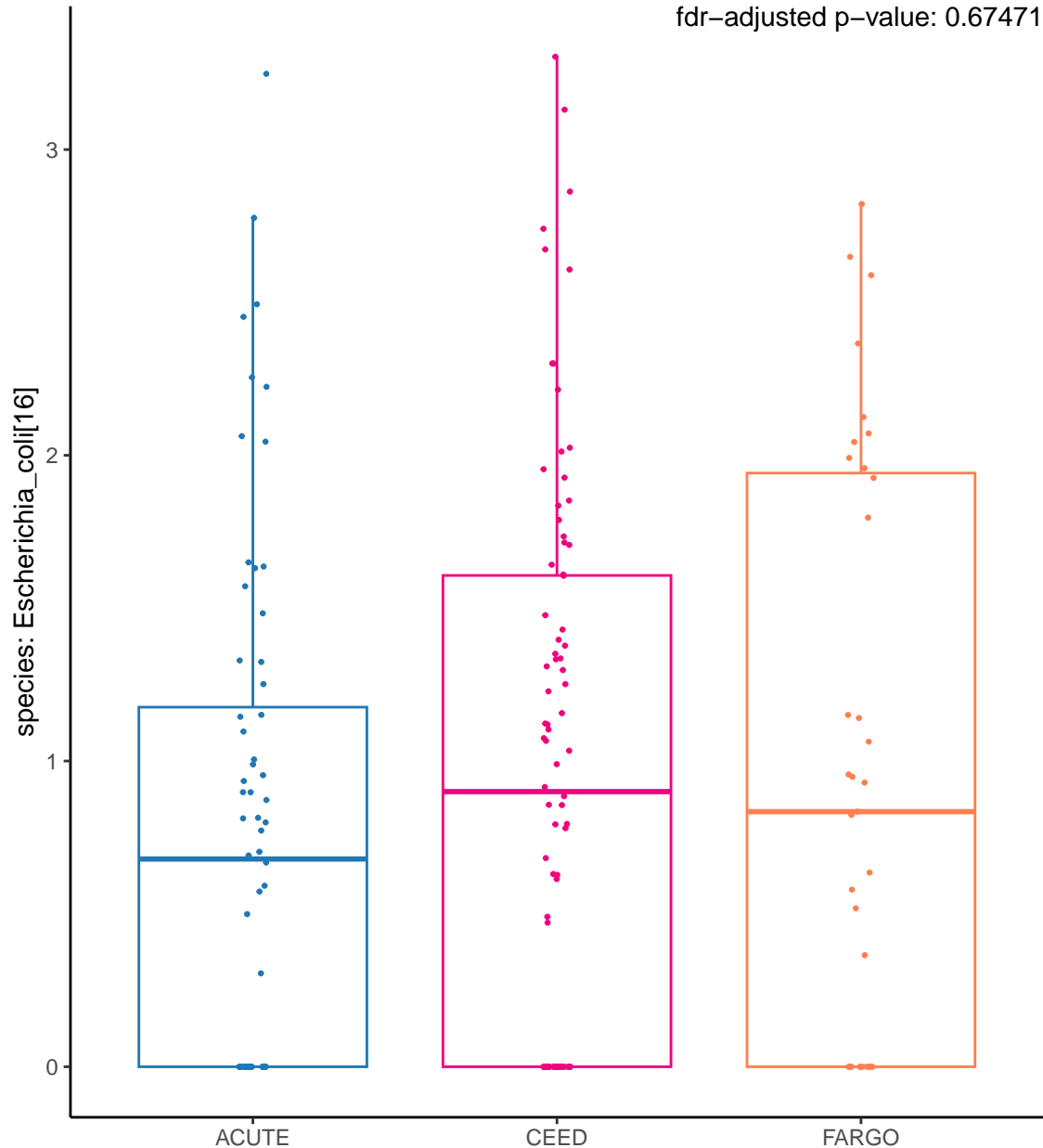
FARGO

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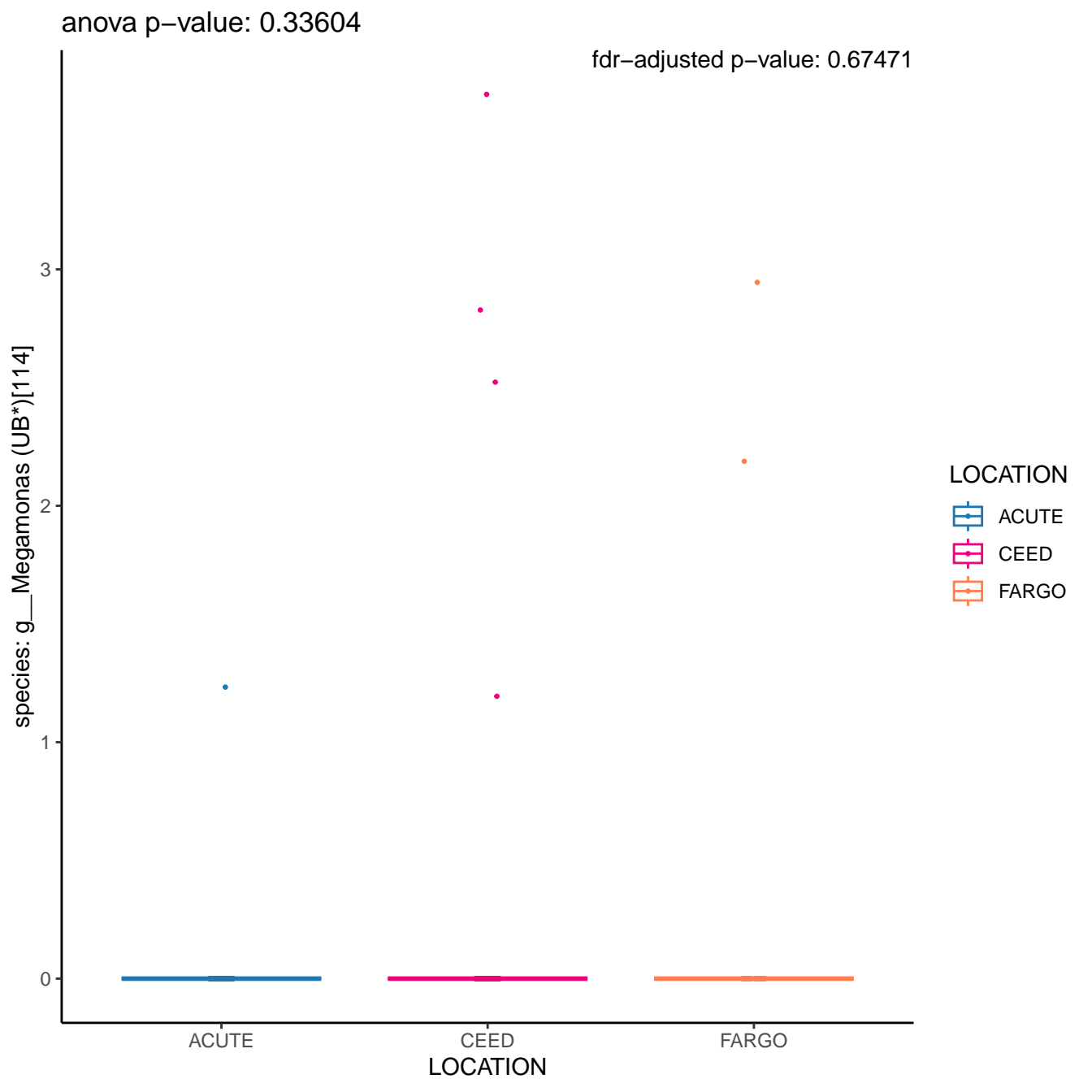
LOCATION



anova p-value: 0.33604

fdr-adjusted p-value: 0.67471

species: g__Megamonas (UB*)[114]



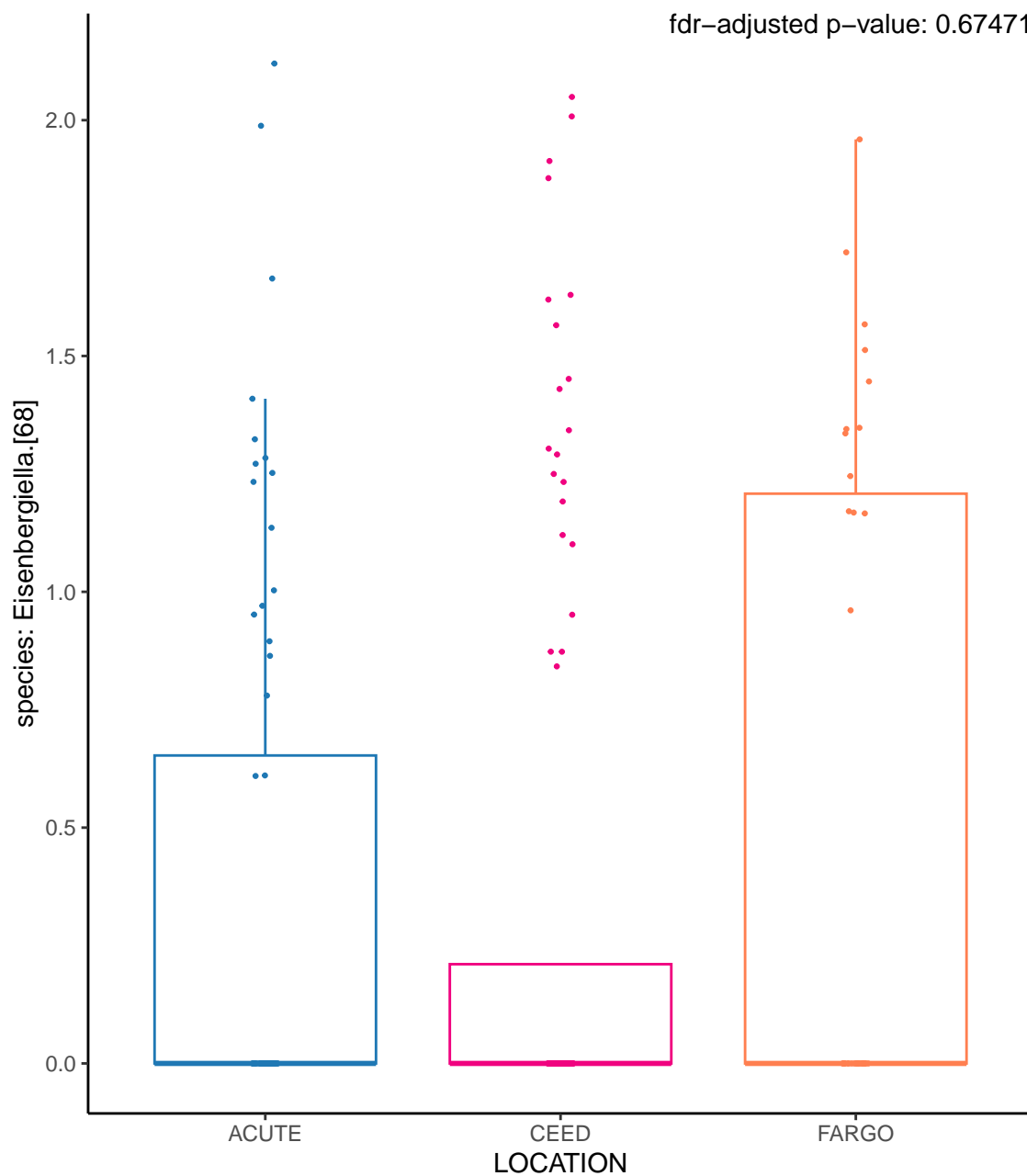
anova p-value: 0.33846

fdr-adjusted p-value: 0.67471

species: Eisenbergiella.[68]

LOCATION

ACUTE
CEED
FARGO



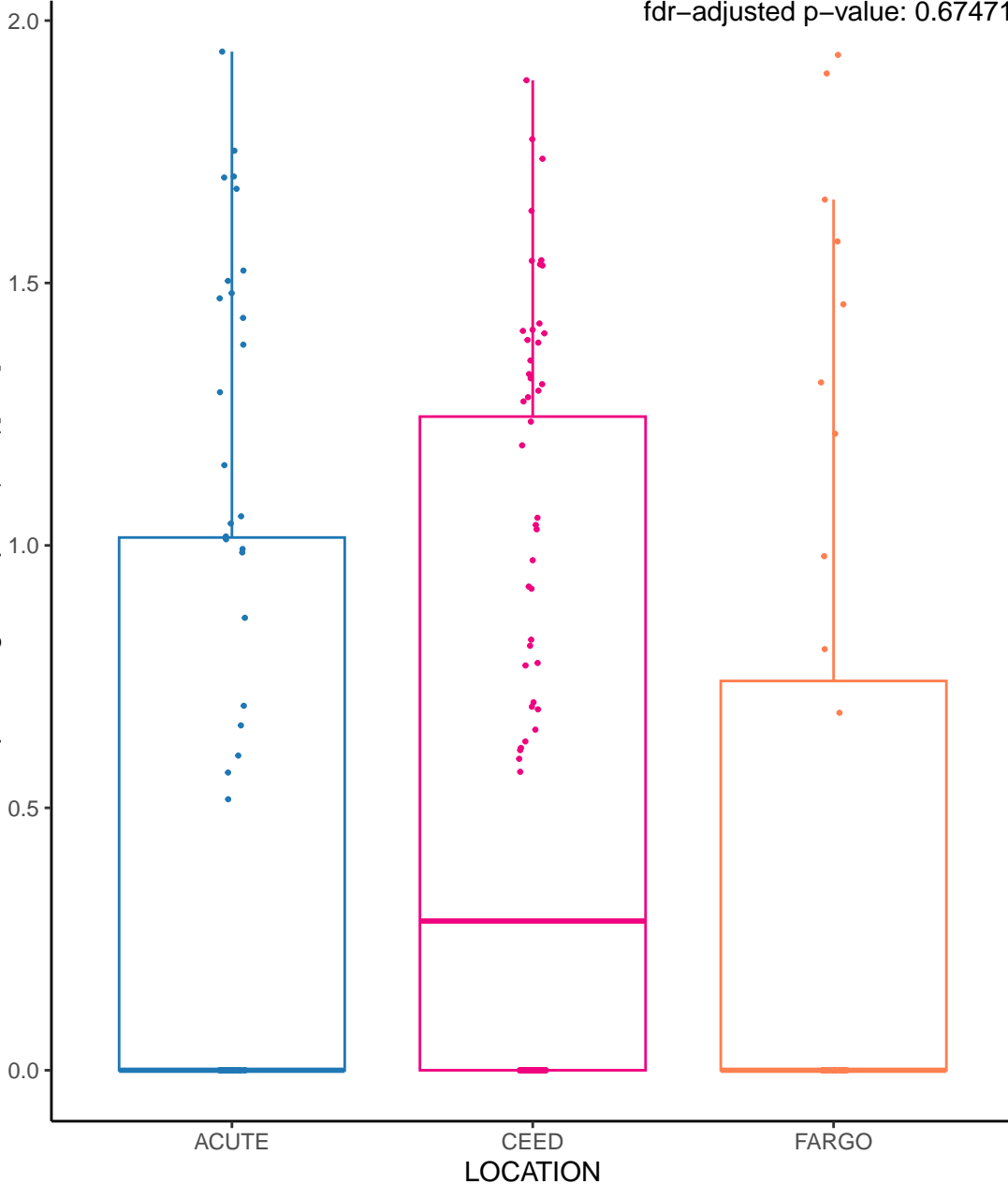
anova p-value: 0.33899

fdr-adjusted p-value: 0.67471

species: g__Bilophila (UB*)[135]

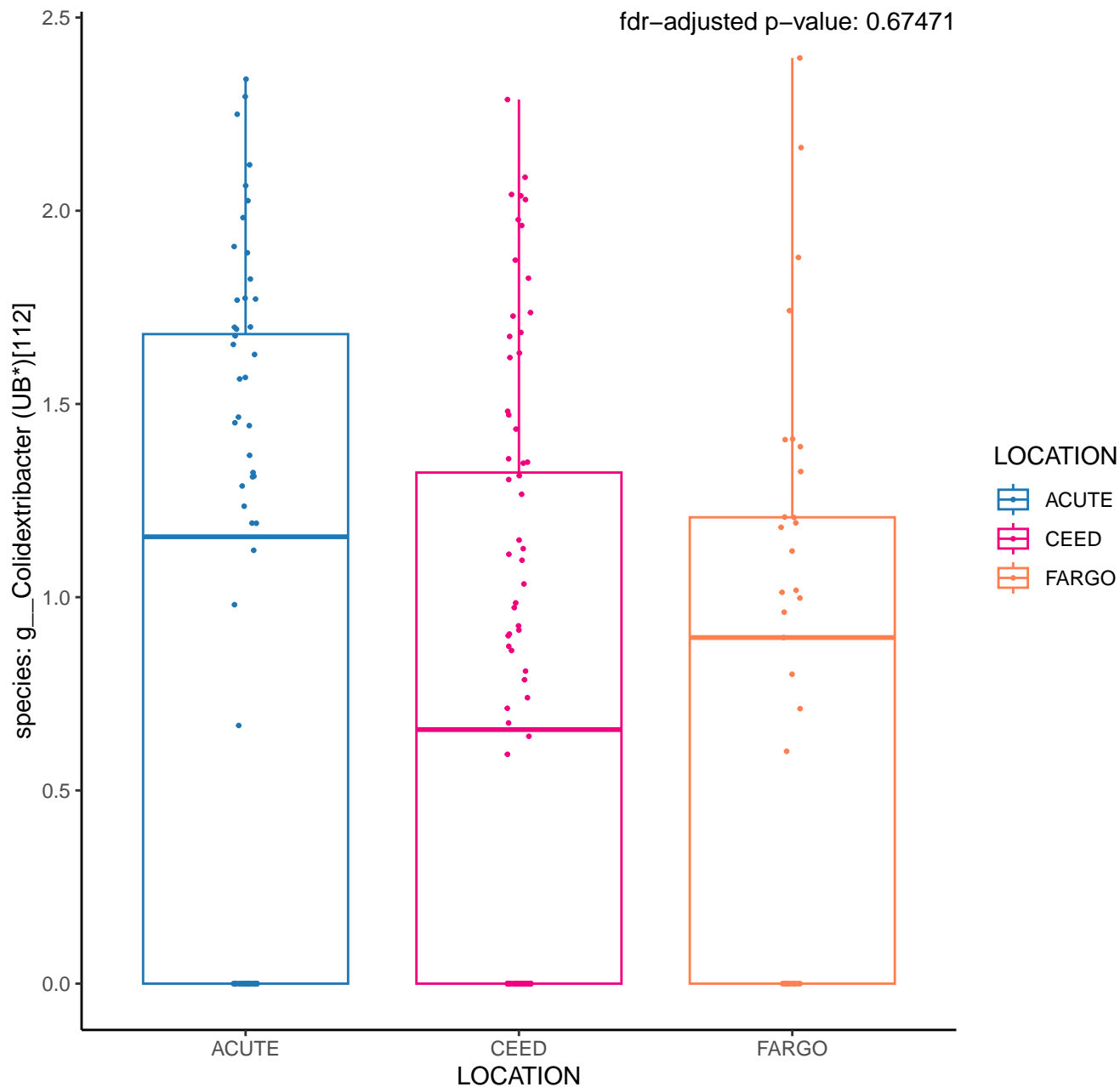
LOCATION

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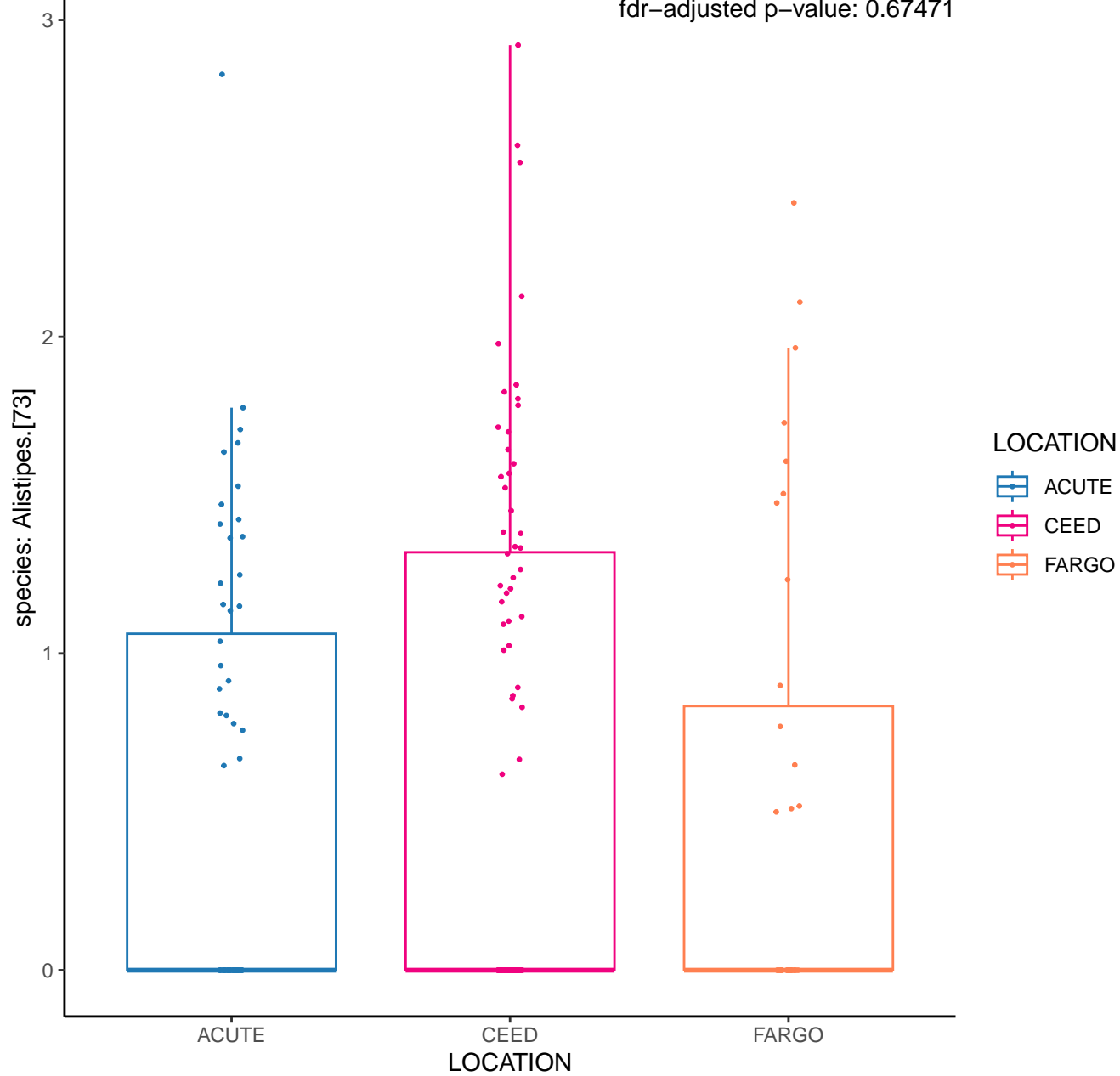
anova p-value: 0.34711

fdr-adjusted p-value: 0.67471



anova p-value: 0.34804

fdr-adjusted p-value: 0.67471



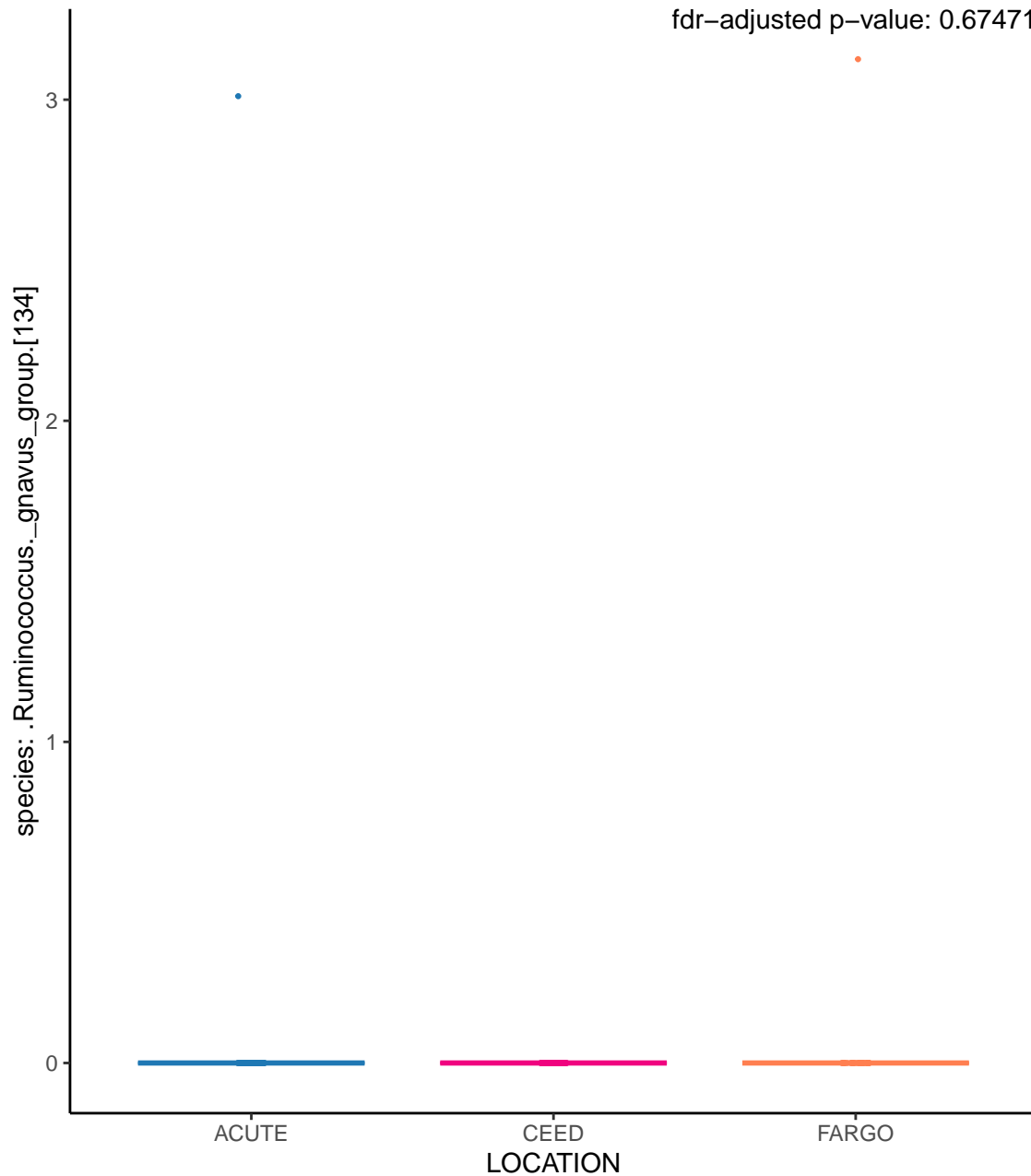
anova p-value: 0.35175

fdr-adjusted p-value: 0.67471

species: .Ruminococcus._gnavus_group.[134]

LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.35304

fdr-adjusted p-value: 0.67471

species: Eggerthellaceae.[122]

LOCATION

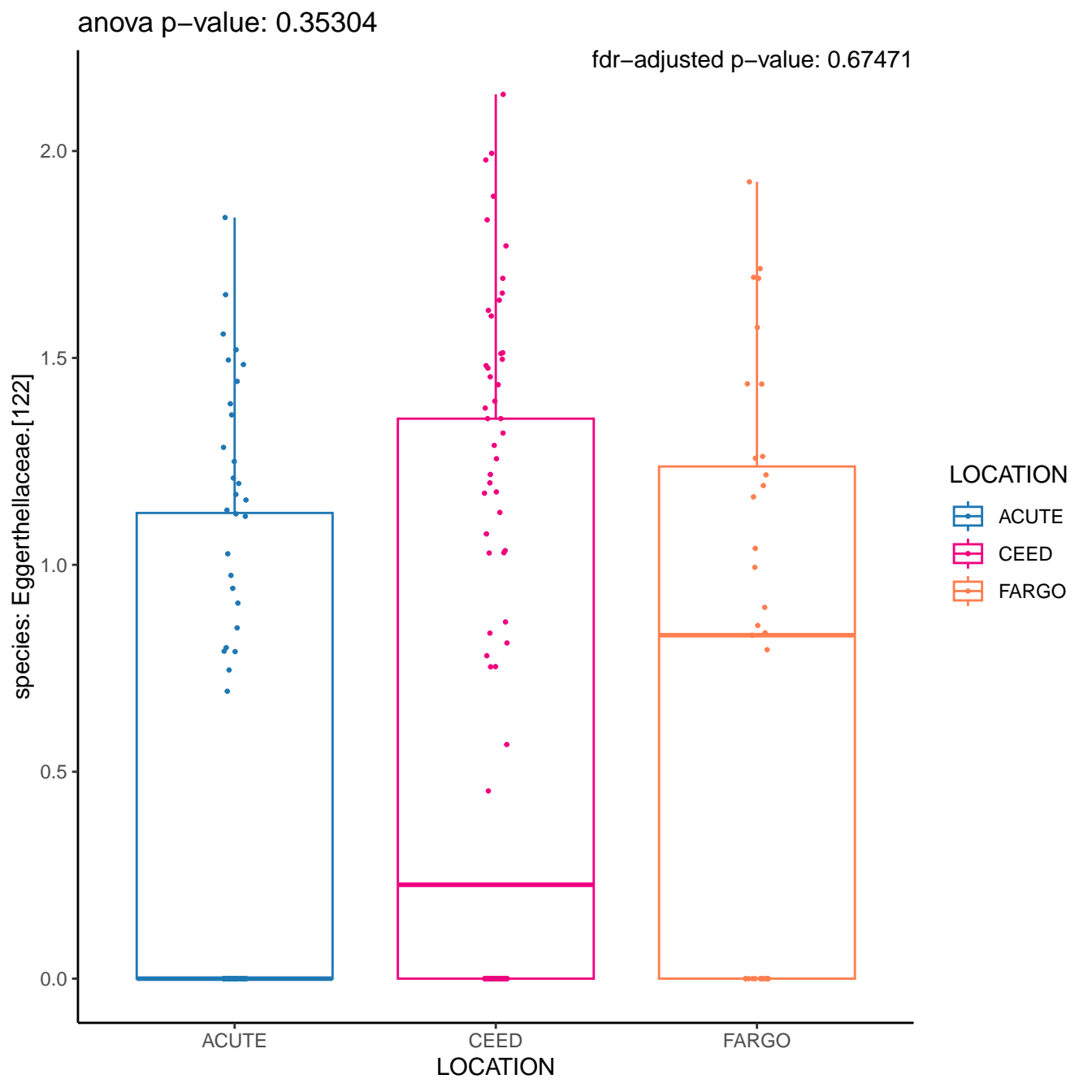
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LOCATION



anova p-value: 0.37168

fdr-adjusted p-value: 0.69225

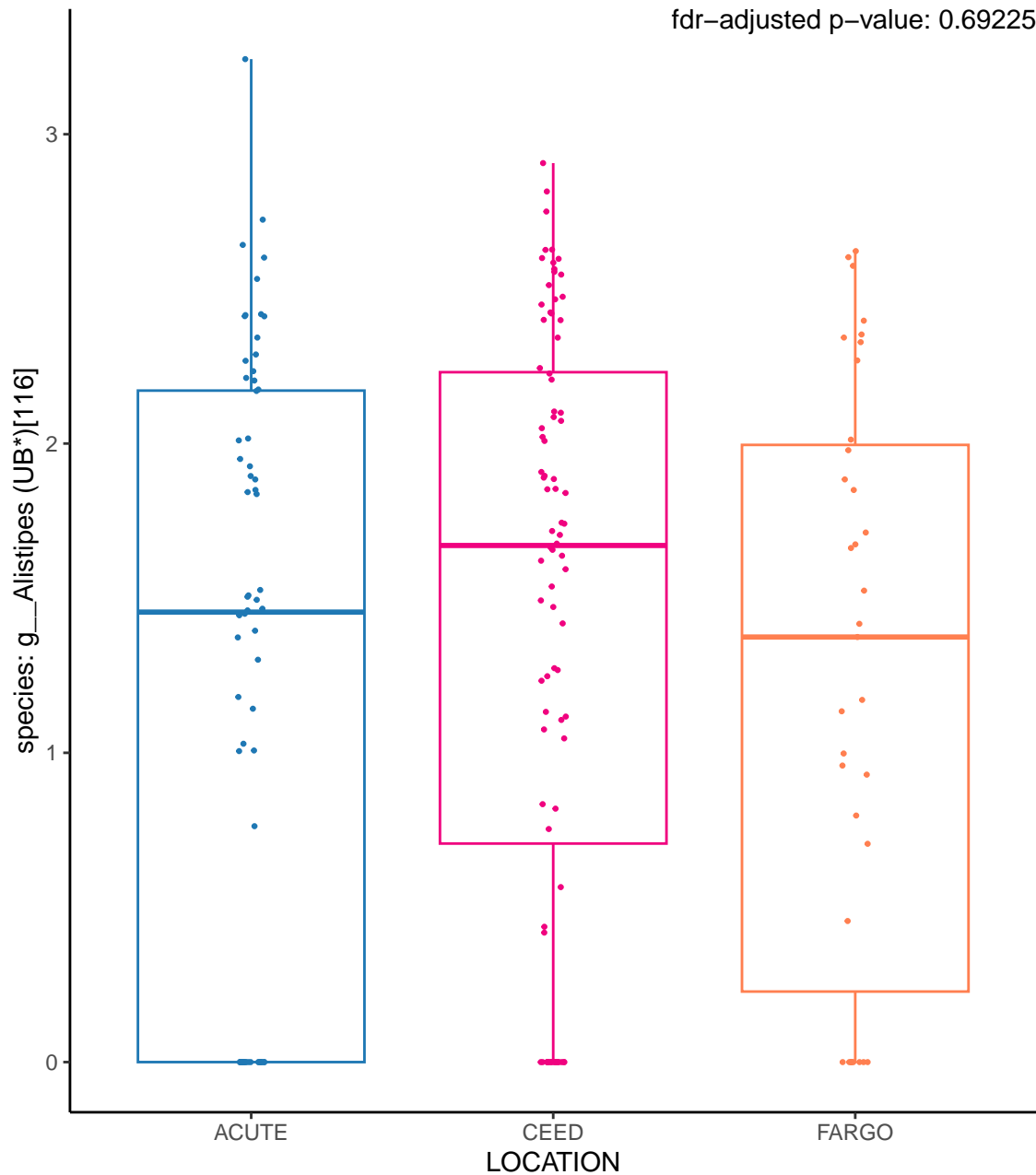
species: g__Alistipes (UB*)[16]

LOCATION

ACUTE

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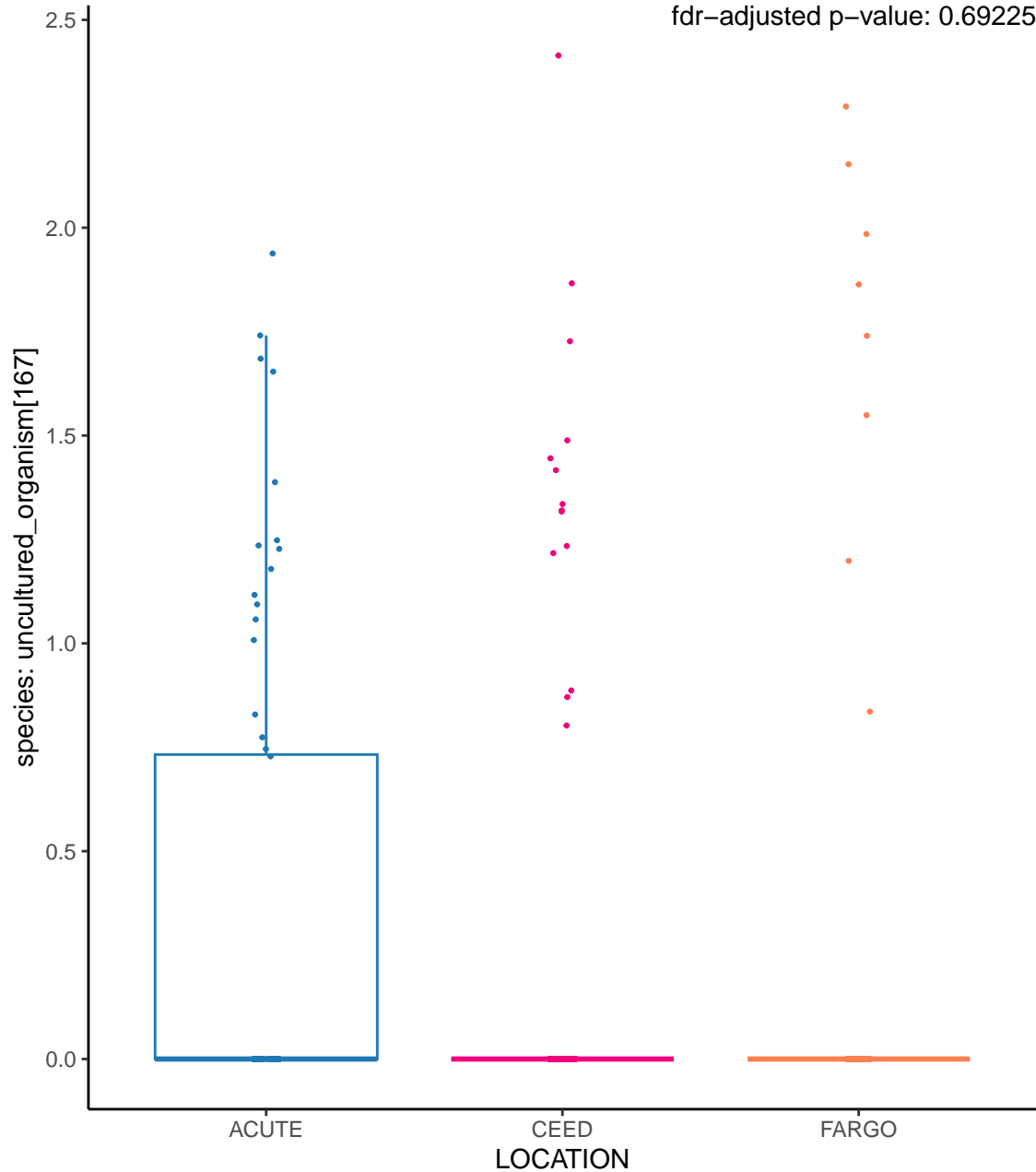
anova p-value: 0.37477

fdr-adjusted p-value: 0.69225

species: uncultured_organism[167]

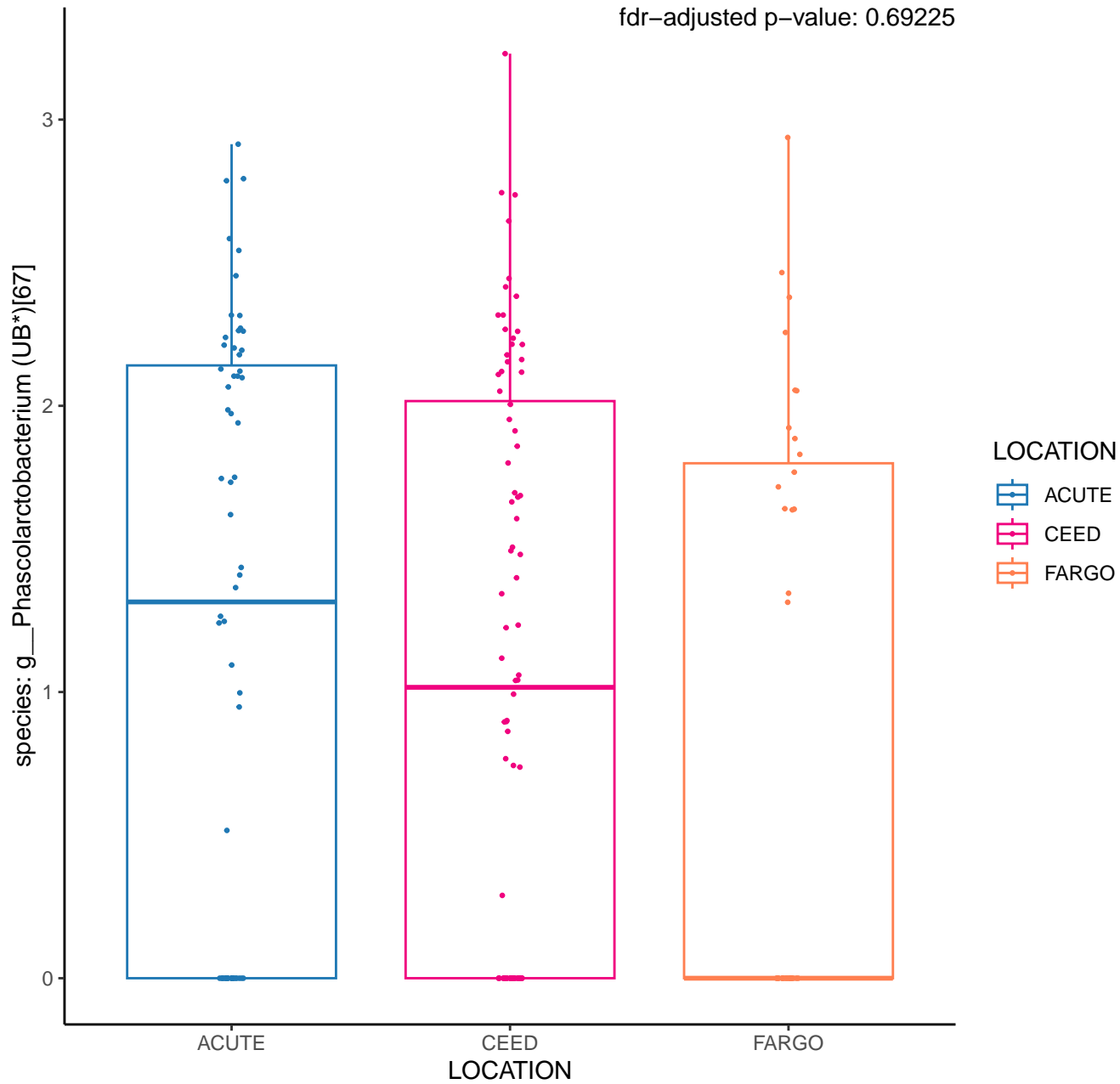
LOCATION

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anova p-value: 0.37489

fdr-adjusted p-value: 0.69225



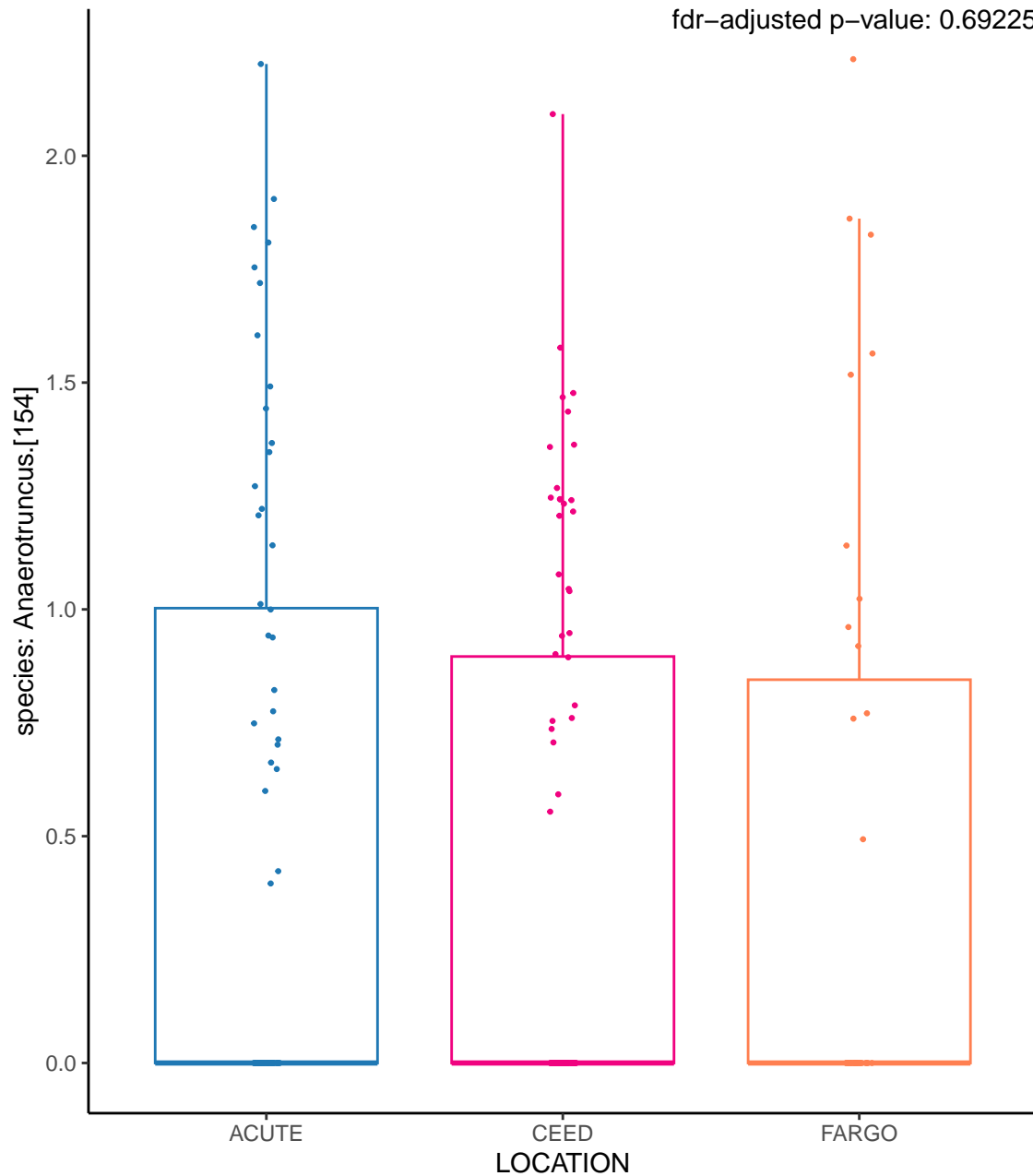
anova p-value: 0.39591

fdr-adjusted p-value: 0.69225

species: Anaerotruncus.[154]

LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.39685

fdr-adjusted p-value: 0.69225

species: *Phascolarctobacterium*. [130]

LOCATION

ACUTE

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FARGO

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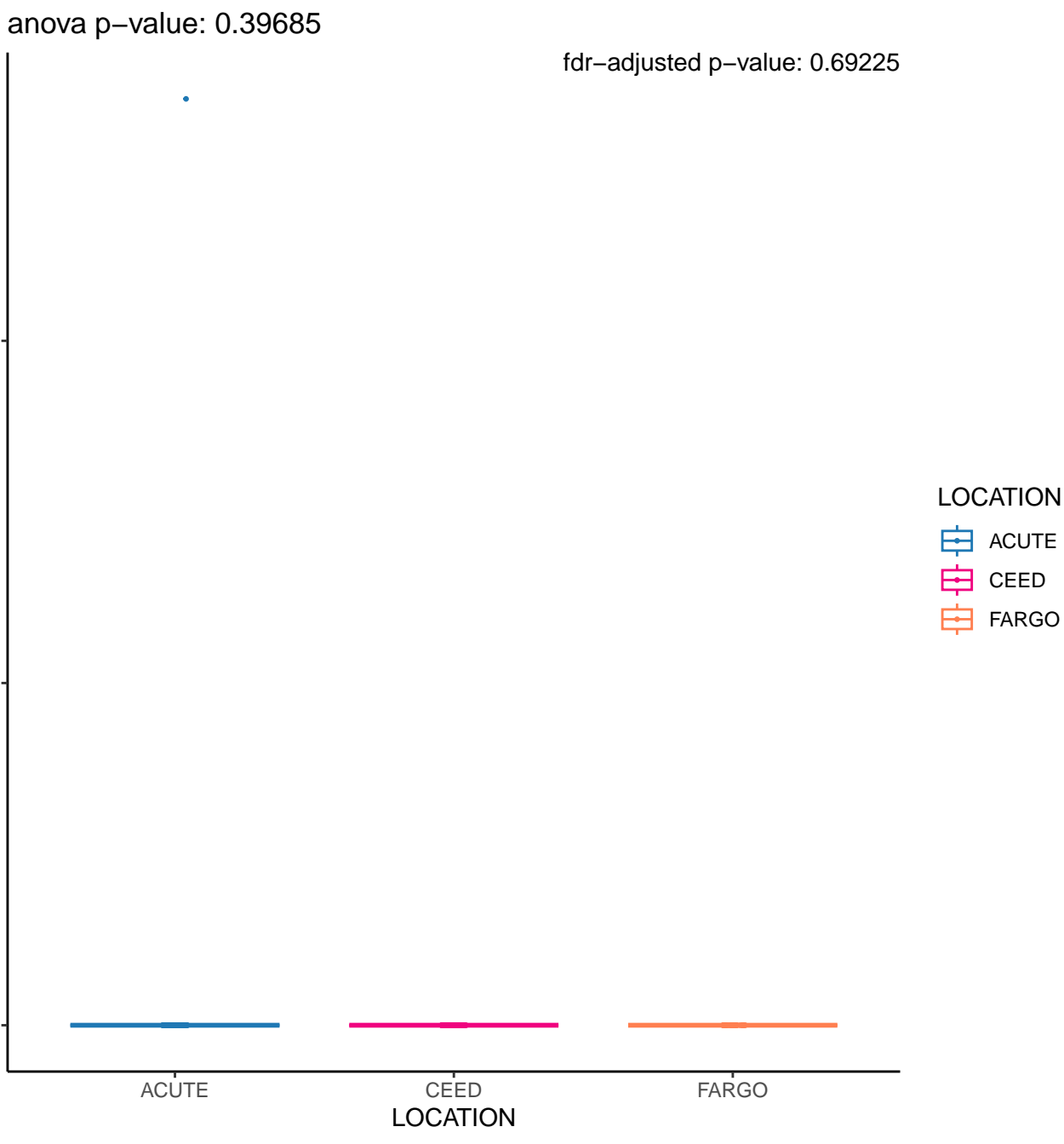
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ACUTE

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FARGO

LOCATION



anova p-value: 0.39685

fdr-adjusted p-value: 0.69225

species: *Dysgonomonas*. [139]

LOCATION

ACUTE

CEED

FARGO

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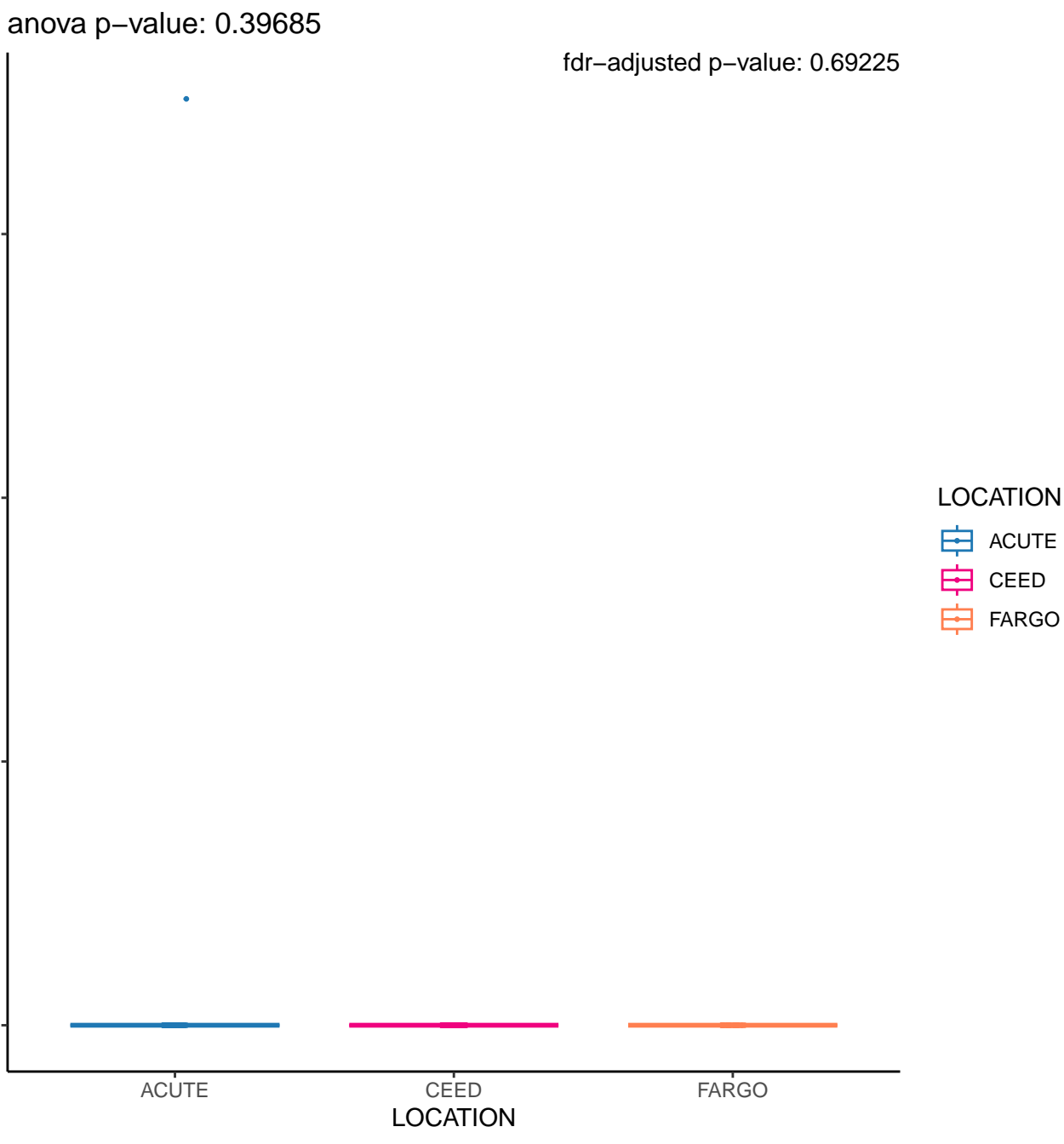
0

ACUTE

CEED

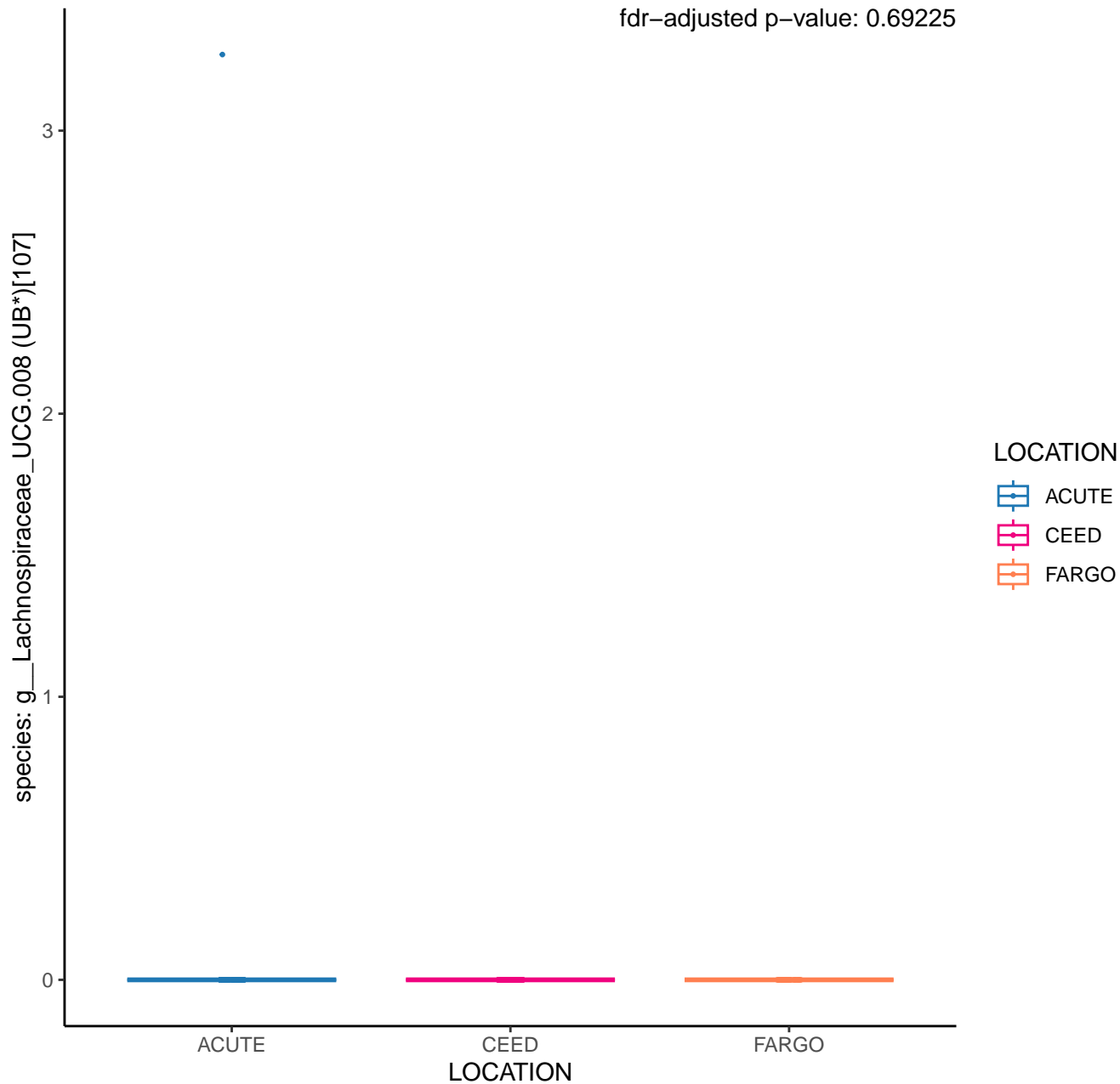
FARGO

LOCATION



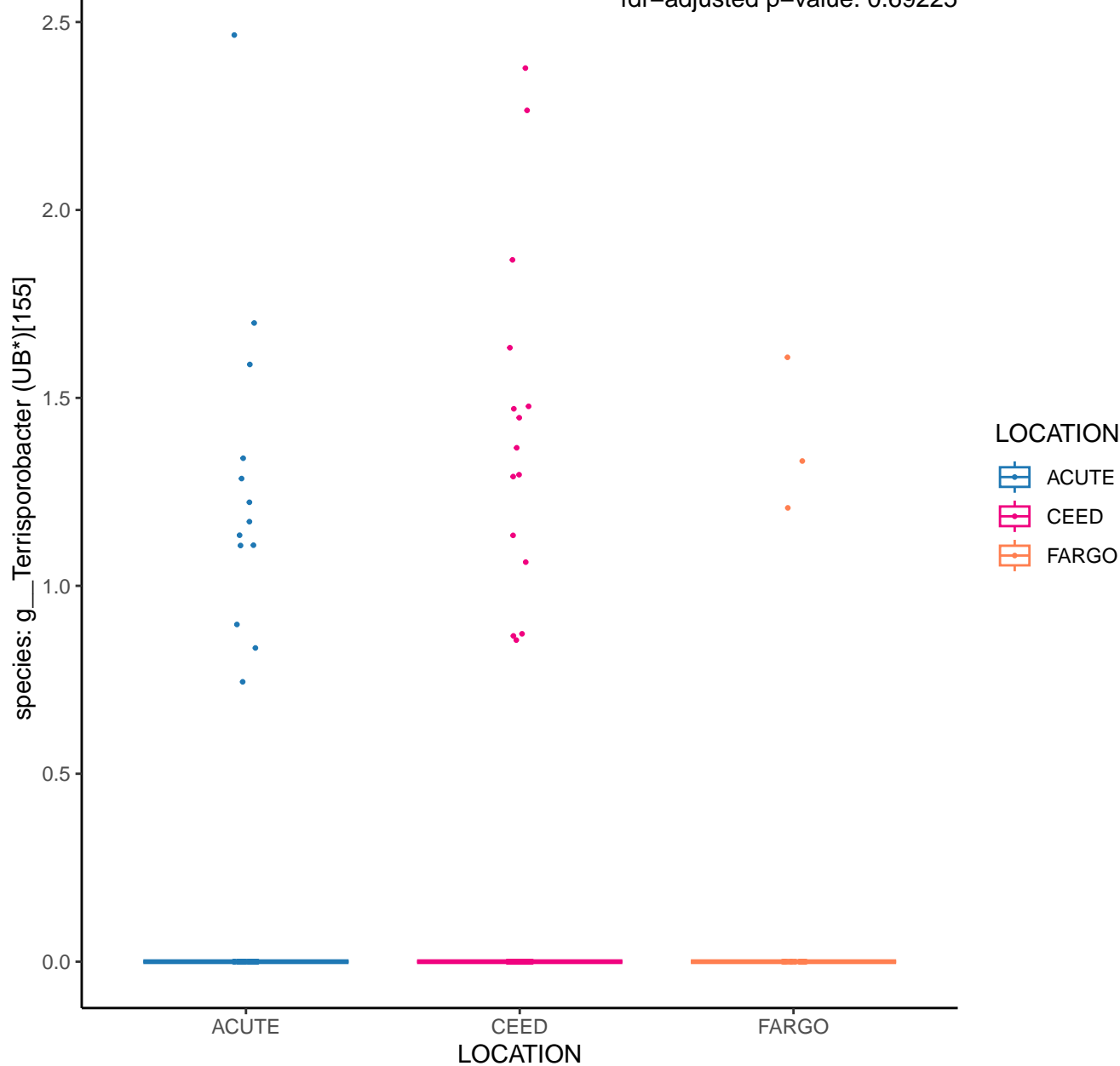
anova p-value: 0.39685

fdr-adjusted p-value: 0.69225



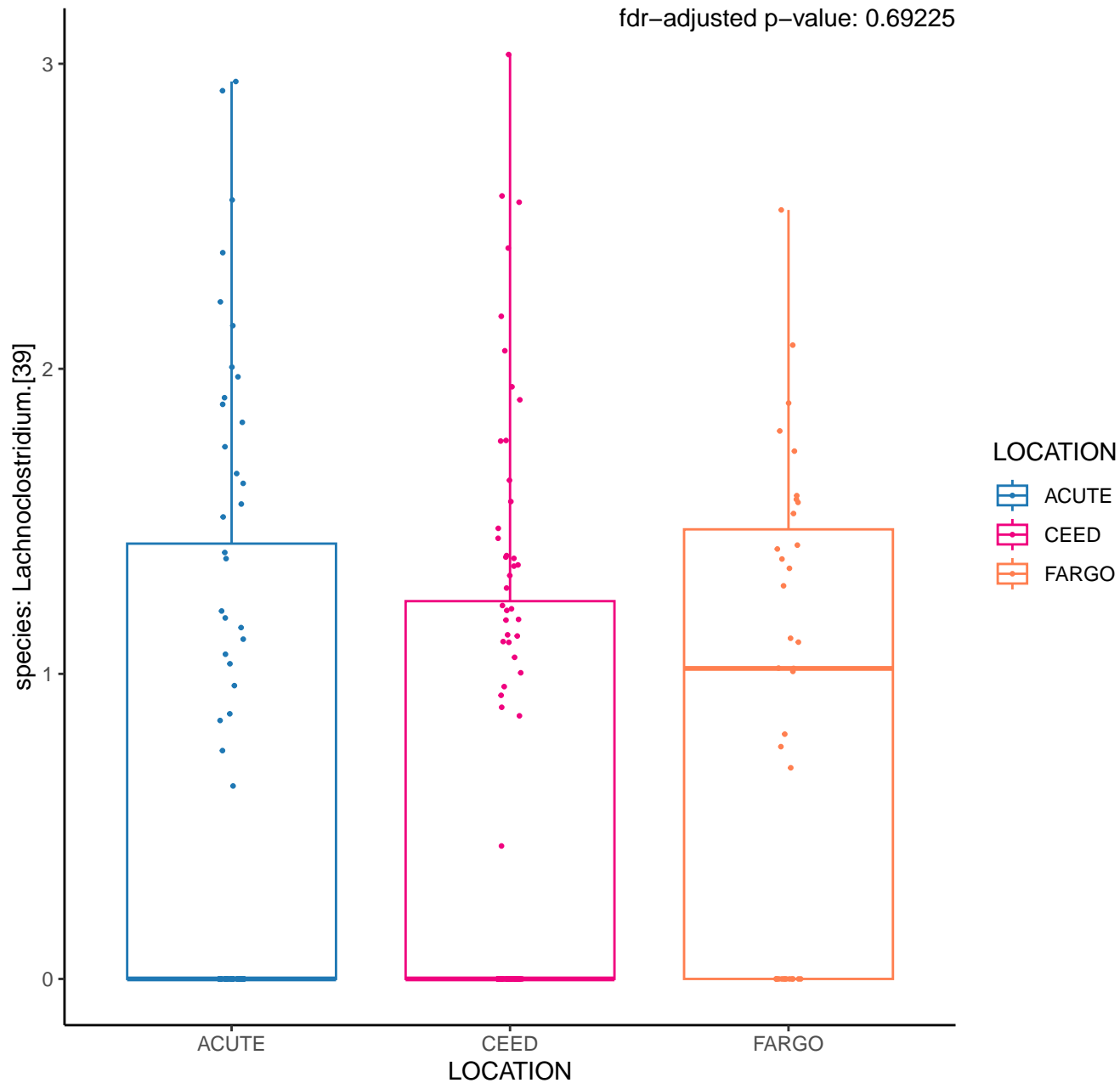
anova p-value: 0.40058

fdr-adjusted p-value: 0.69225



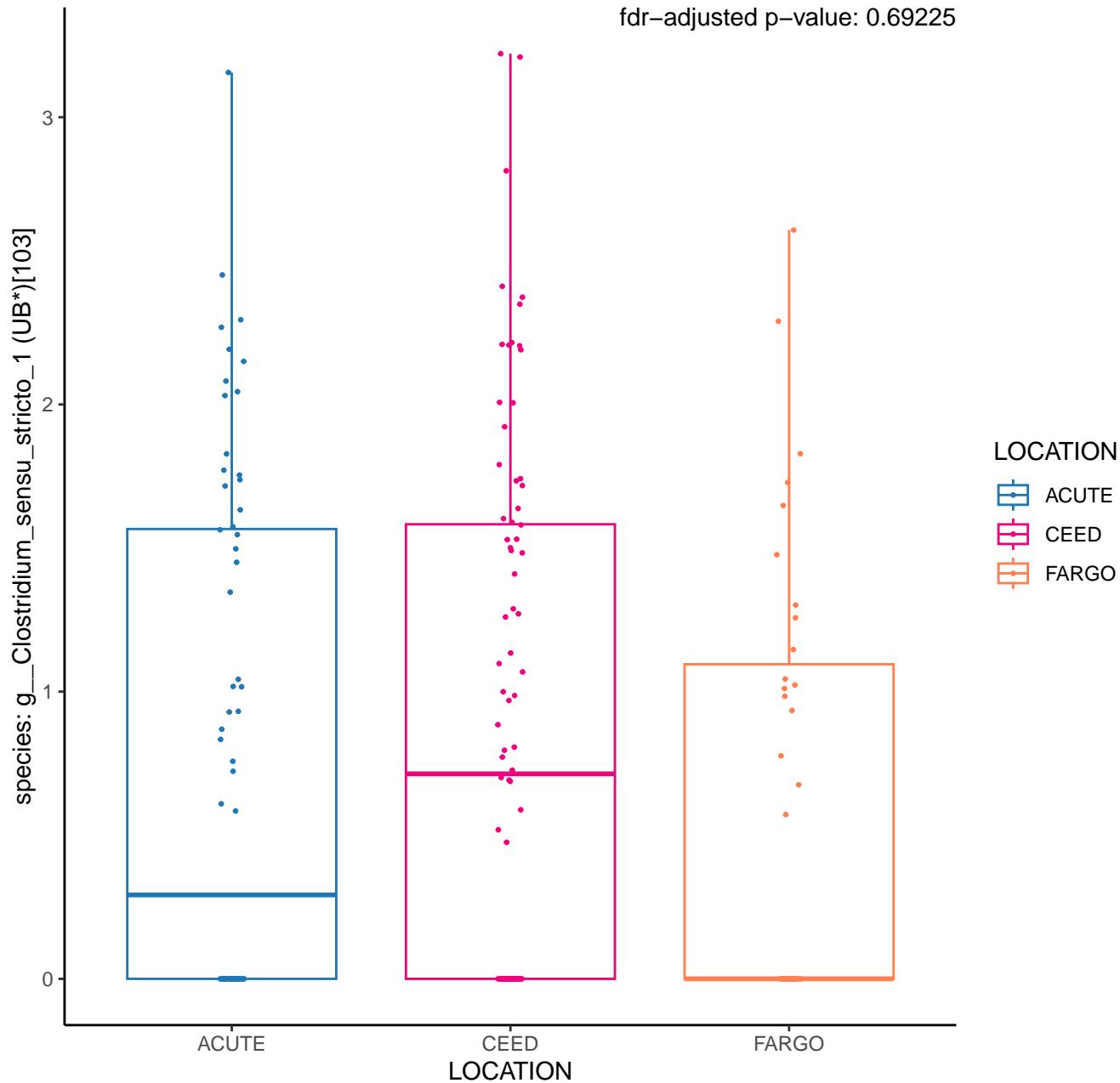
anova p-value: 0.40209

fdr-adjusted p-value: 0.69225



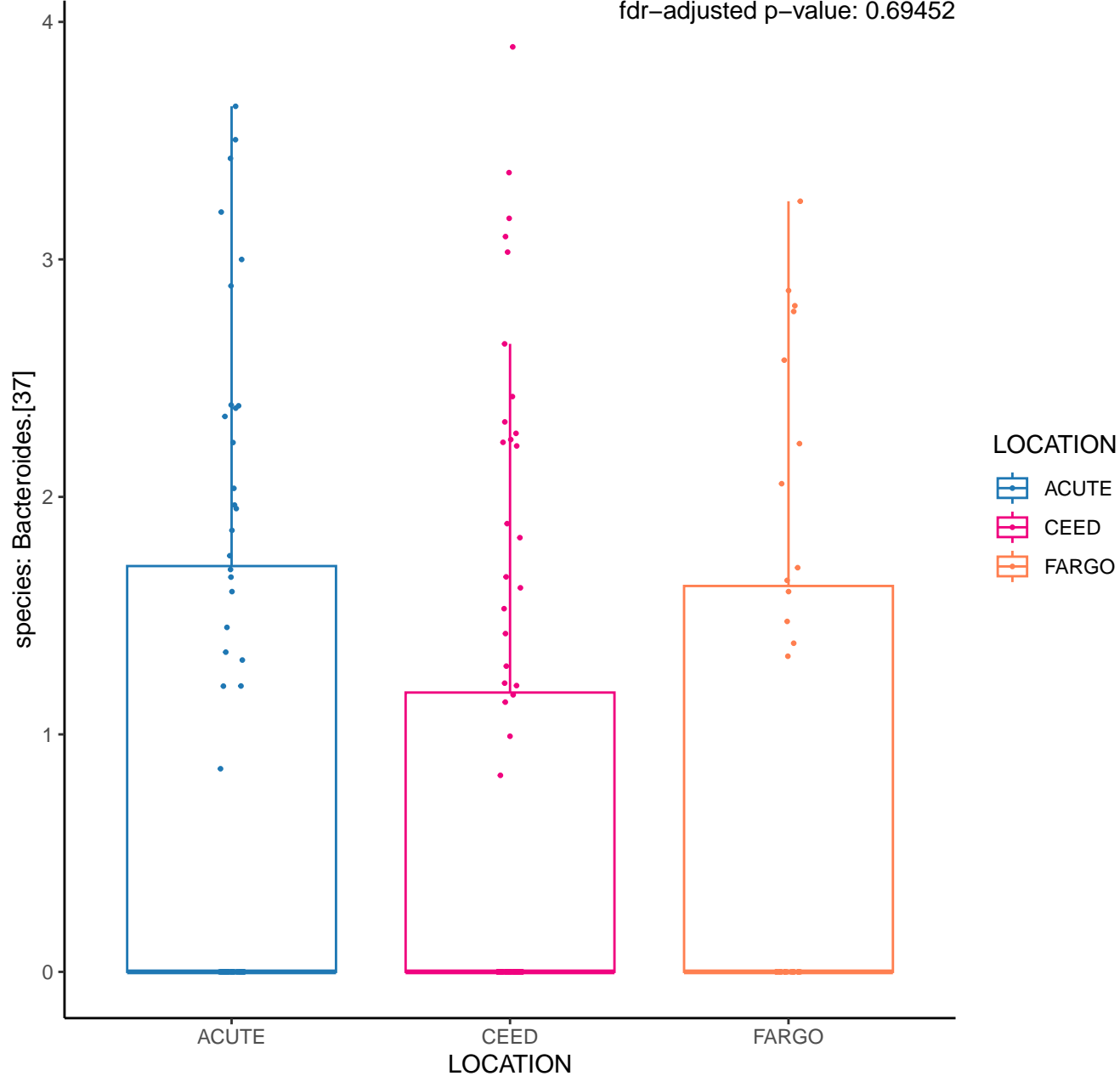
anova p-value: 0.40247

fdr-adjusted p-value: 0.69225



anova p-value: 0.4114

fdr-adjusted p-value: 0.69452



anova p-value: 0.41526

fdr-adjusted p-value: 0.69452

species: g__uncultured (UB*)[124]

LOCATION

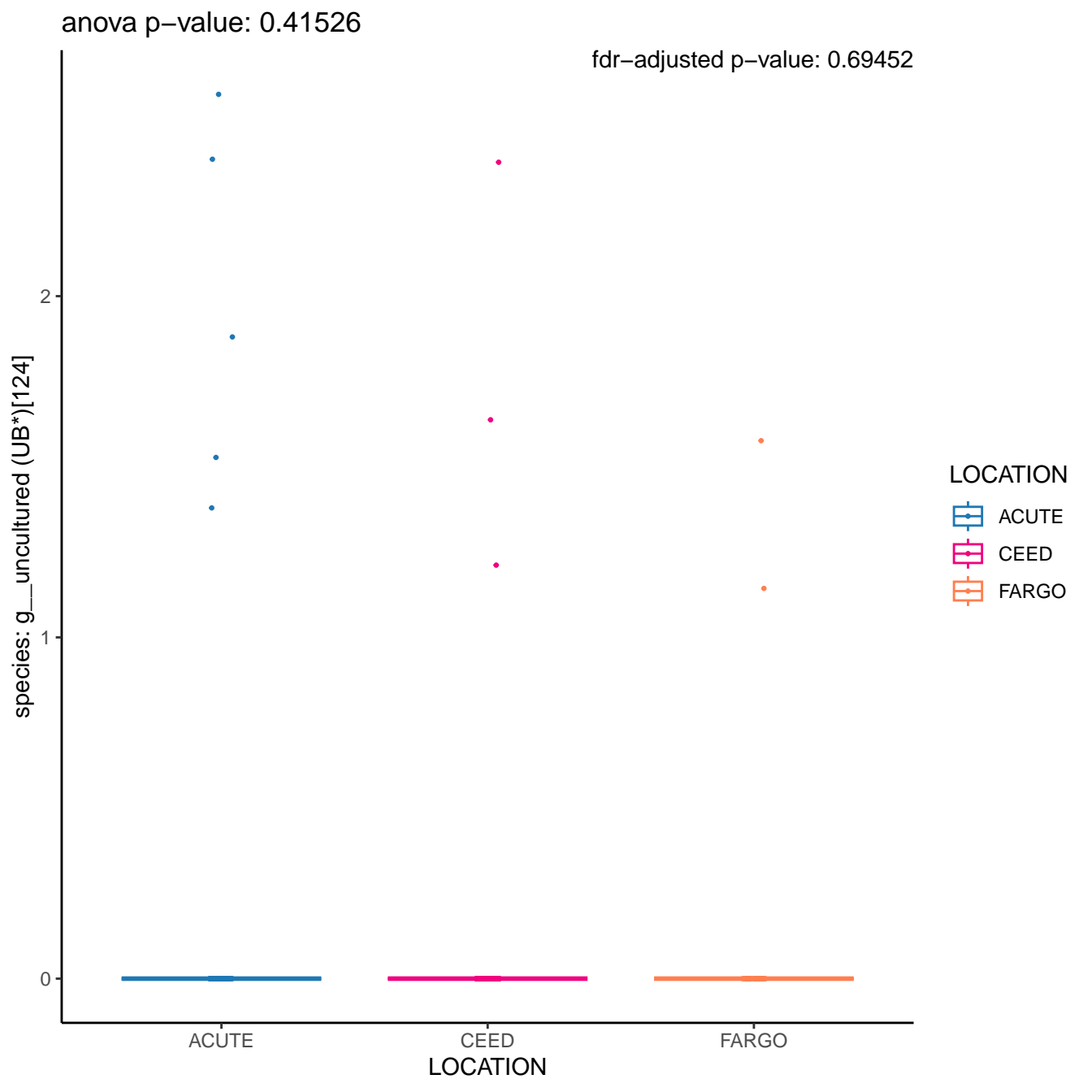
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LOCATION



anova p-value: 0.42055

fdr-adjusted p-value: 0.69452

species: g__uncultured (UB*)[159]

LOCATION

ACUTE

CEED

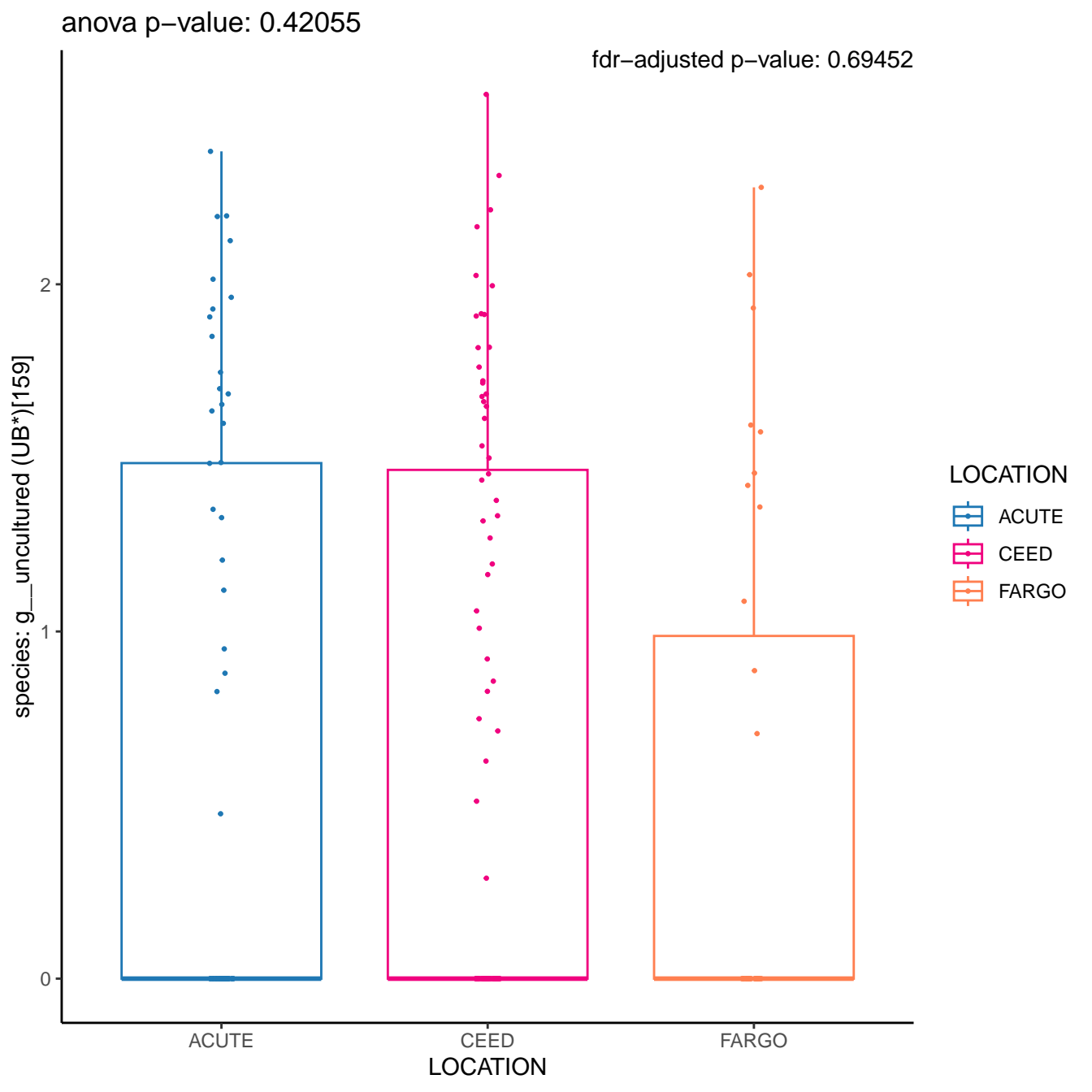
FARGO

ACUTE

CEED

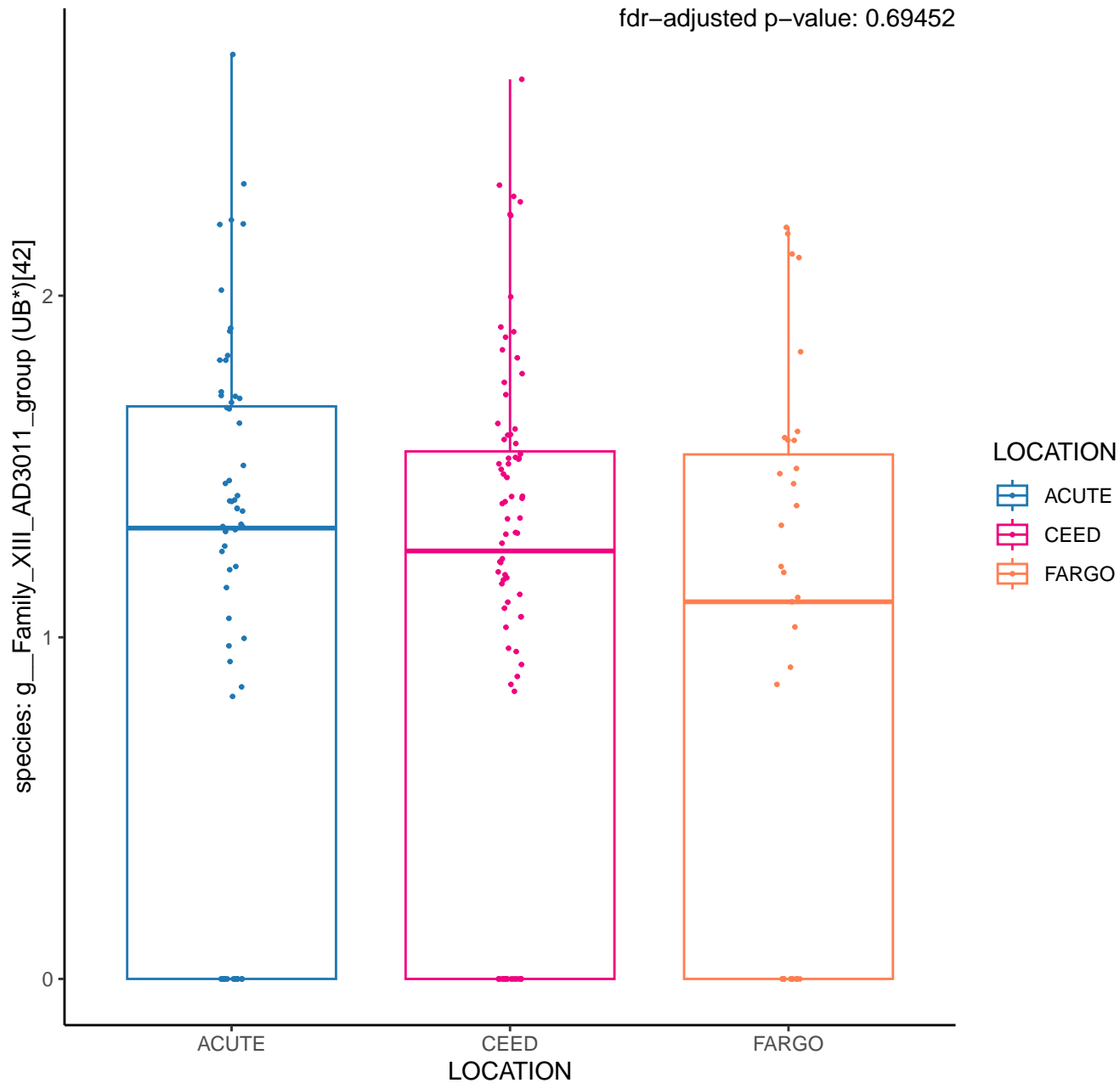
FARGO

LOCATION



anova p-value: 0.42496

fdr-adjusted p-value: 0.69452



anova p-value: 0.42683

fdr-adjusted p-value: 0.69452

species: Methanobrevibacter. [69]

LOCATION

ACUTE

CEED

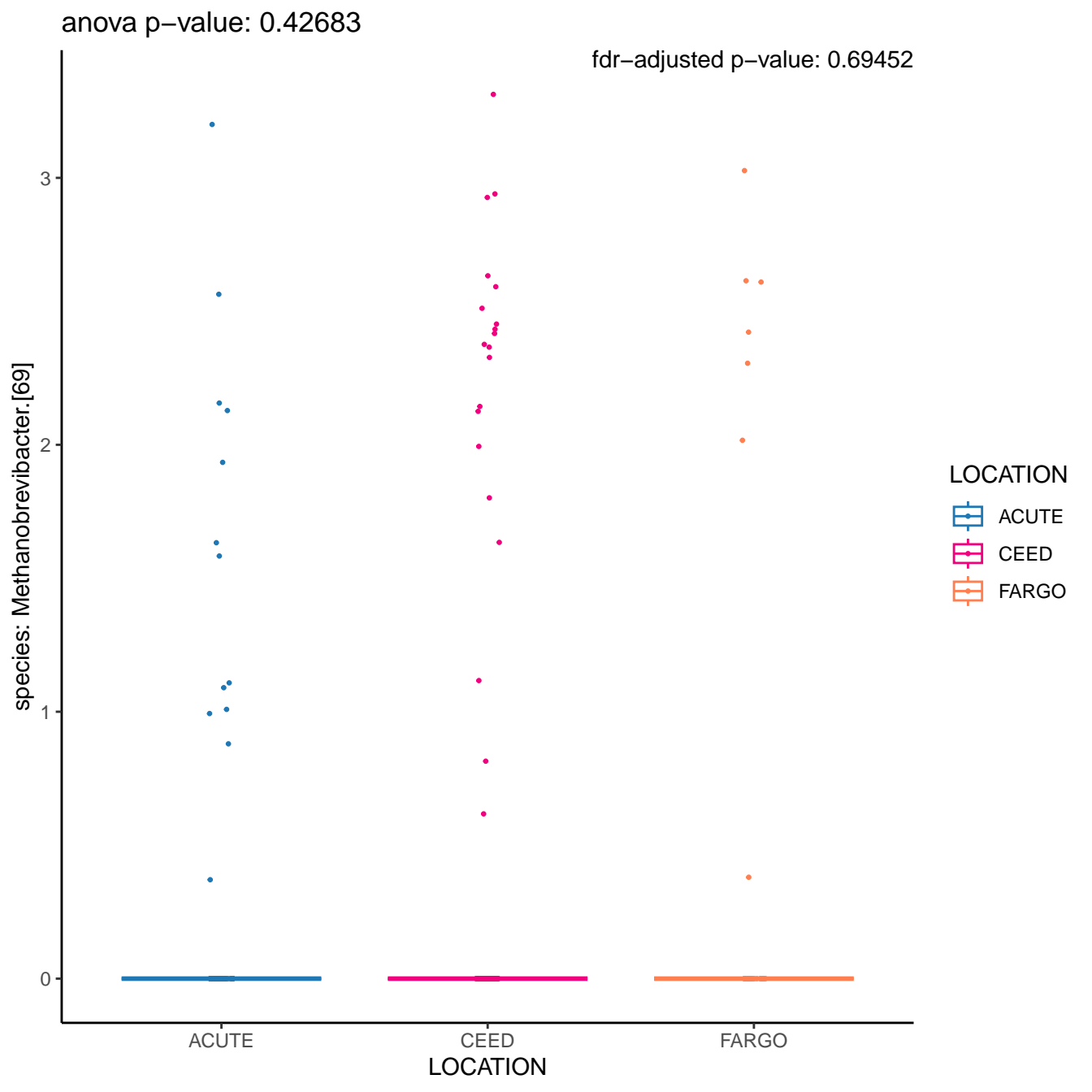
FARGO

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LOCATION



anova p-value: 0.42802

fdr-adjusted p-value: 0.69452

species: *Lactobacillus*. [85]

LOCATION

ACUTE

CEED

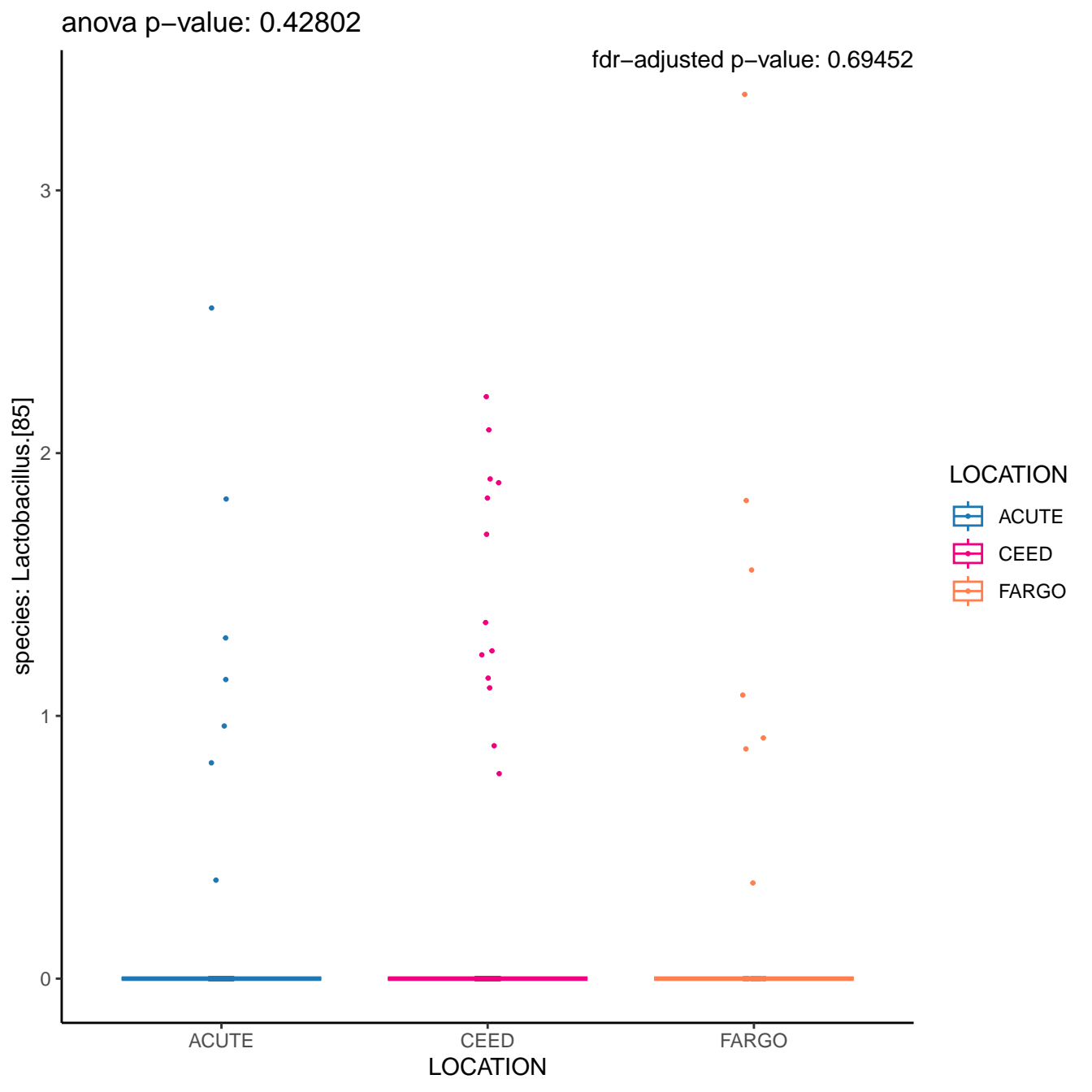
FARGO

ACUTE

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FARGO

LOCATION



anova p-value: 0.4362

fdr-adjusted p-value: 0.69702

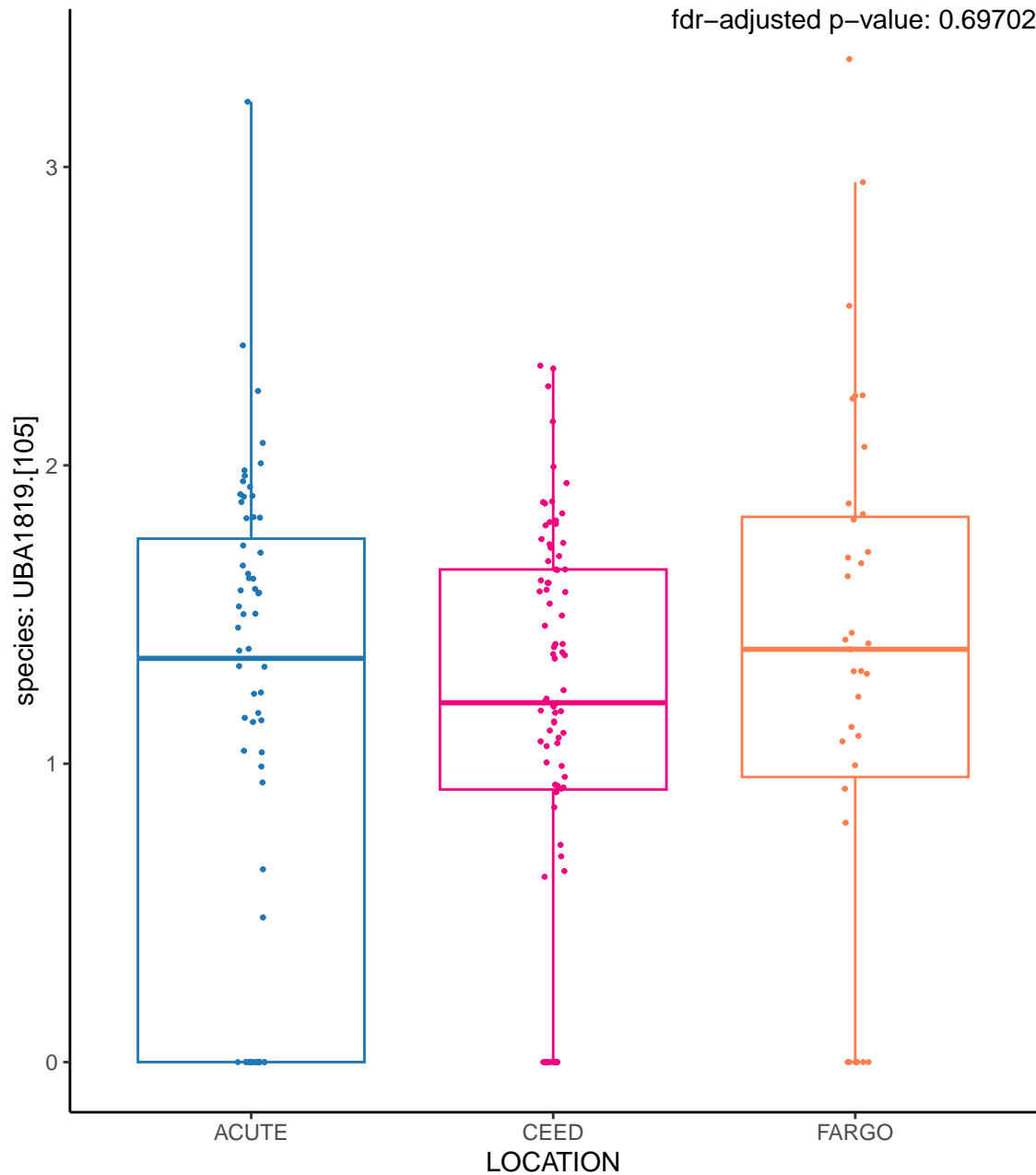
species: UBA1819.[105]

LOCATION

ACUTE

CEED

FARGO



anova p-value: 0.44031

fdr-adjusted p-value: 0.69702

species: g__Shuttleworthia (UB*)[6]

LOCATION

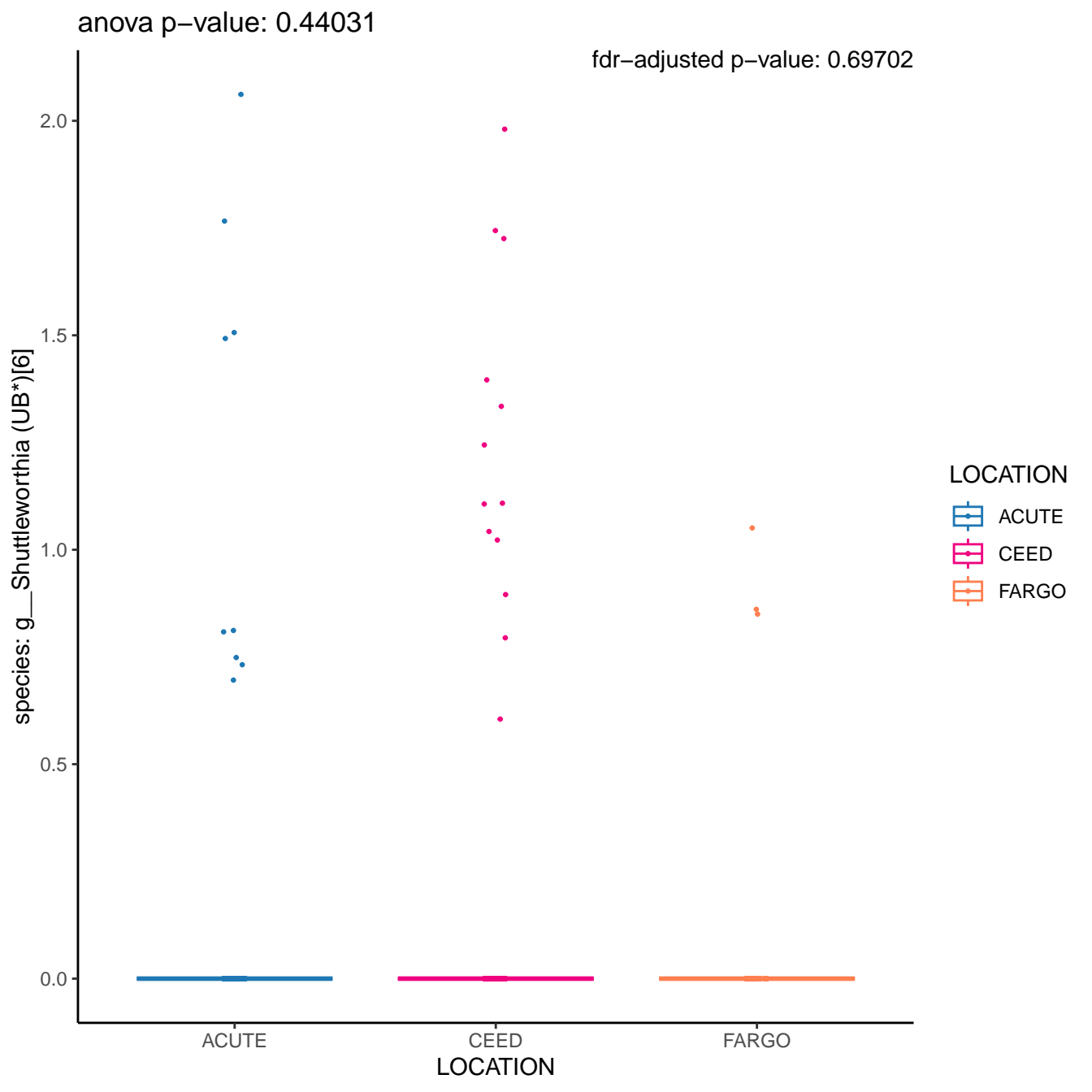
ACUTE
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ACUTE

CEED

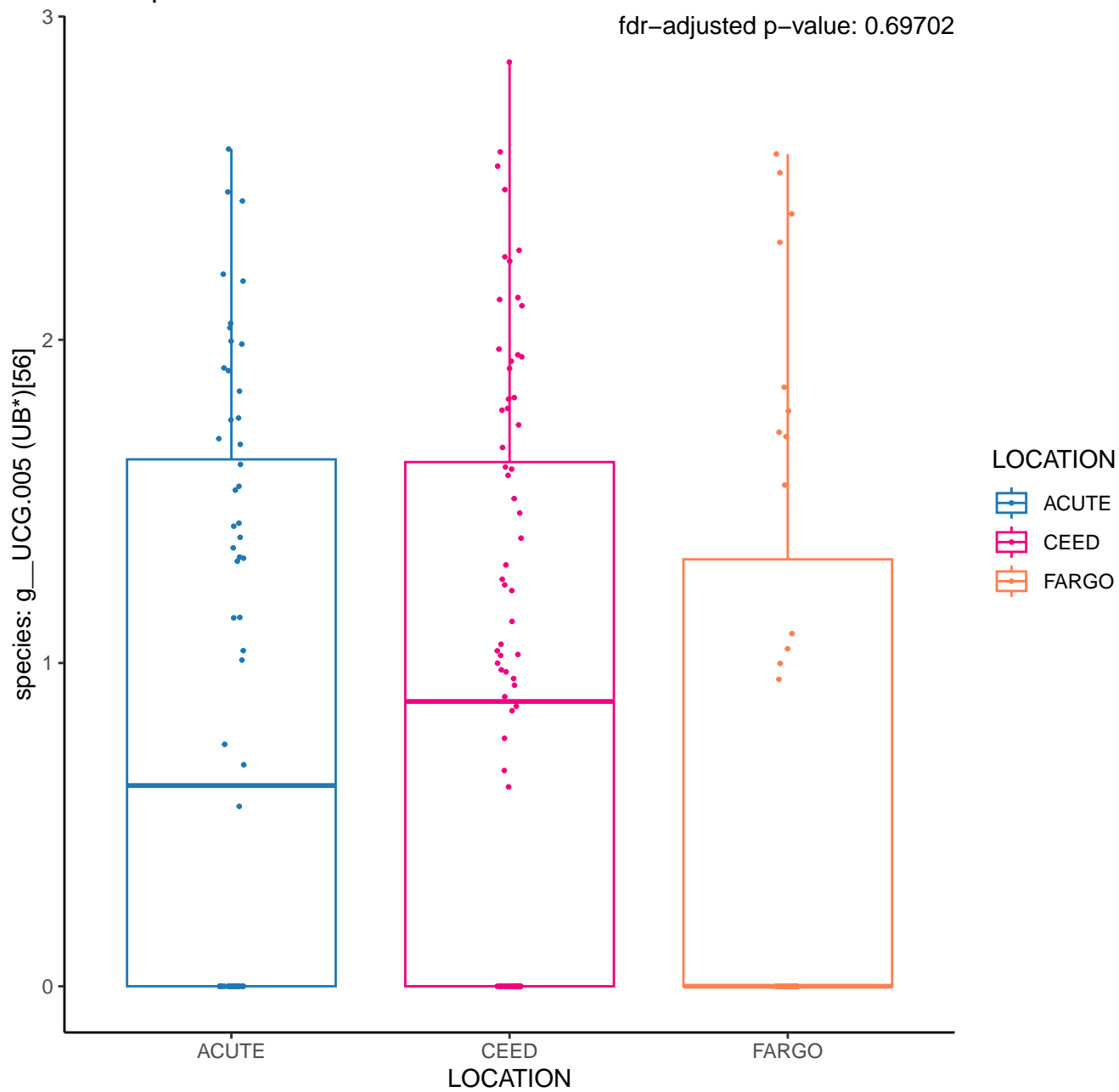
FARGO

LOCATION



anova p-value: 0.44595

fdr-adjusted p-value: 0.69702



anova p-value: 0.44627

fdr-adjusted p-value: 0.69702

species: Erysipelotrichaceae_UCG.003.[132]

LOCATION

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ACUTE

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LOCATION

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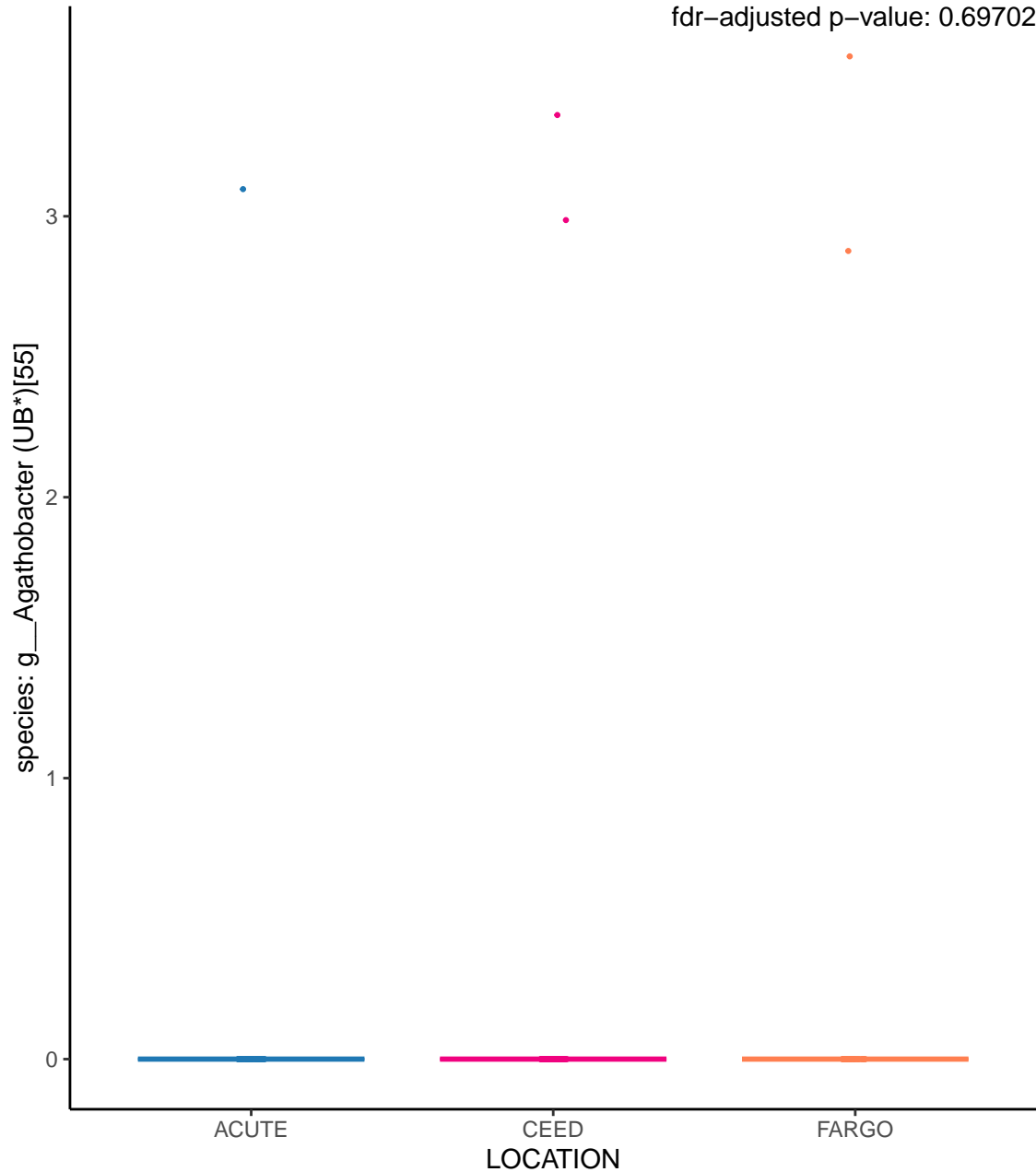
anova p-value: 0.44982

fdr-adjusted p-value: 0.69702

species: g__Agathobacter (UB*)[55]

LOCATION

ACUTE
CEED
FARGO

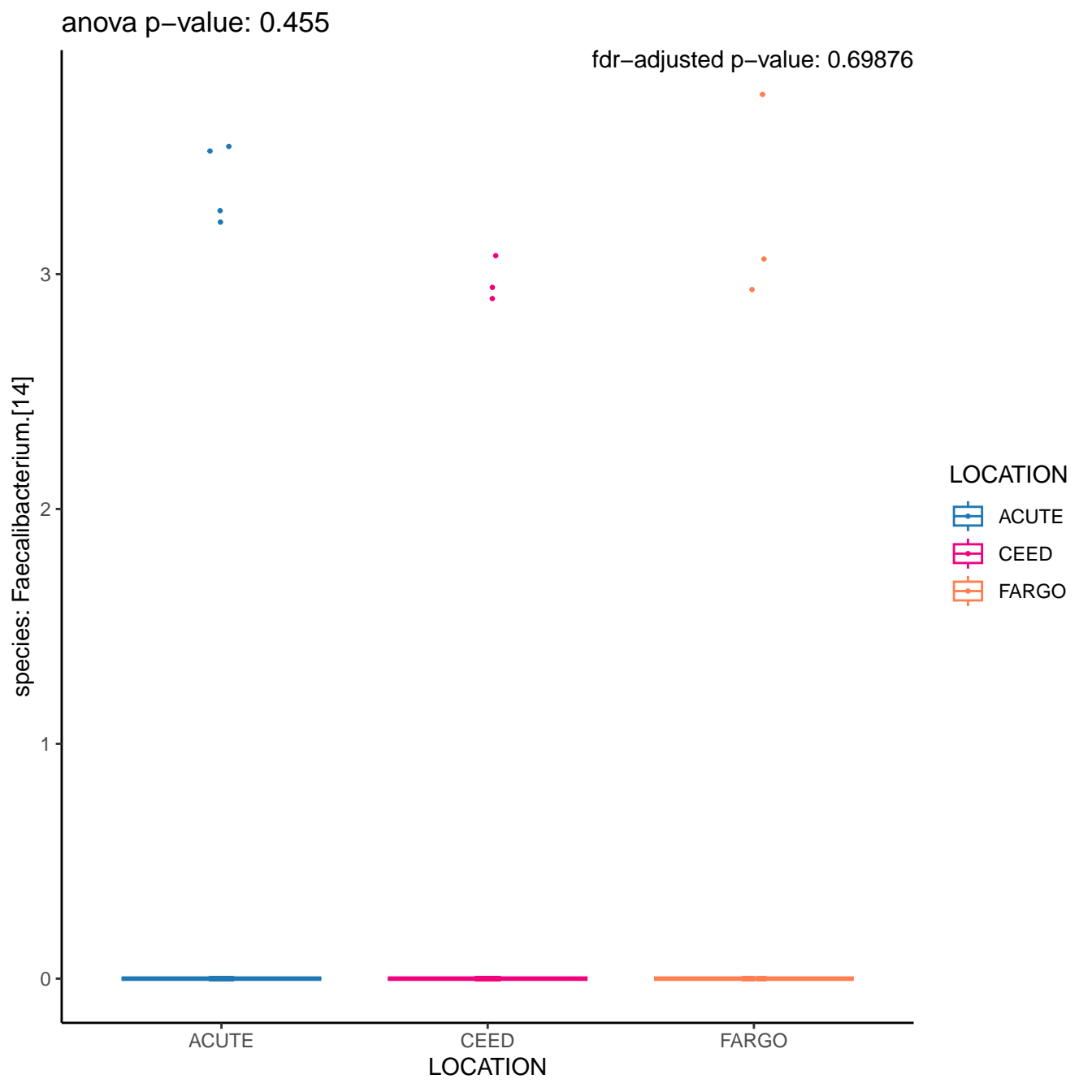


anova p-value: 0.455

fdr-adjusted p-value: 0.69876

species: *Faecalibacterium*. [14]

LOCATION



anova p-value: 0.46951

fdr-adjusted p-value: 0.71466

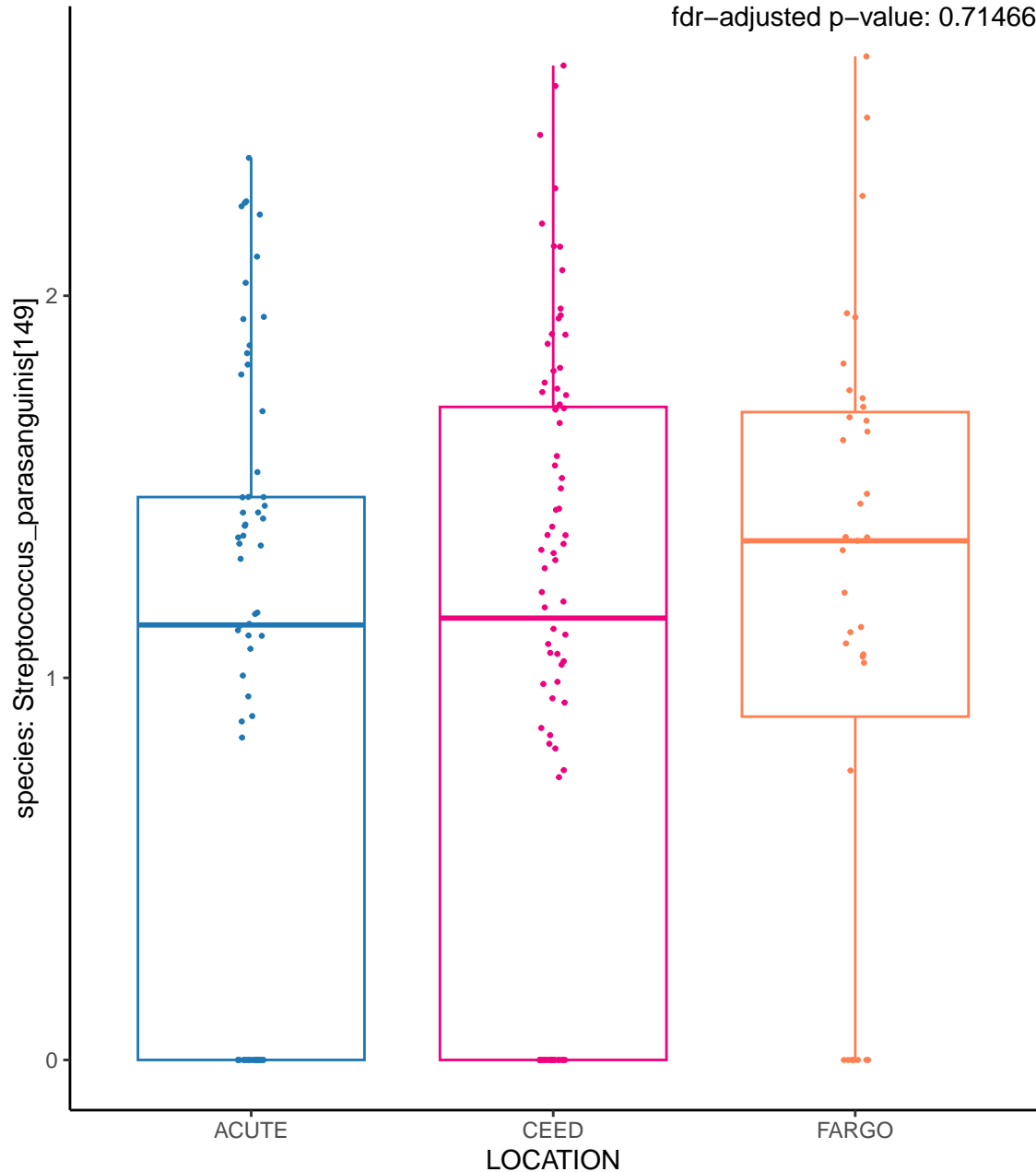
species: *Streptococcus parasanguinis*[149]

LOCATION

ACUTE

CEED

FARGO



anova p-value: 0.49382

fdr-adjusted p-value: 0.74023

species: *Solobacterium*.[143]

LOCATION

ACUTE

CEED

FARGO

species: *Solobacterium*.[143]

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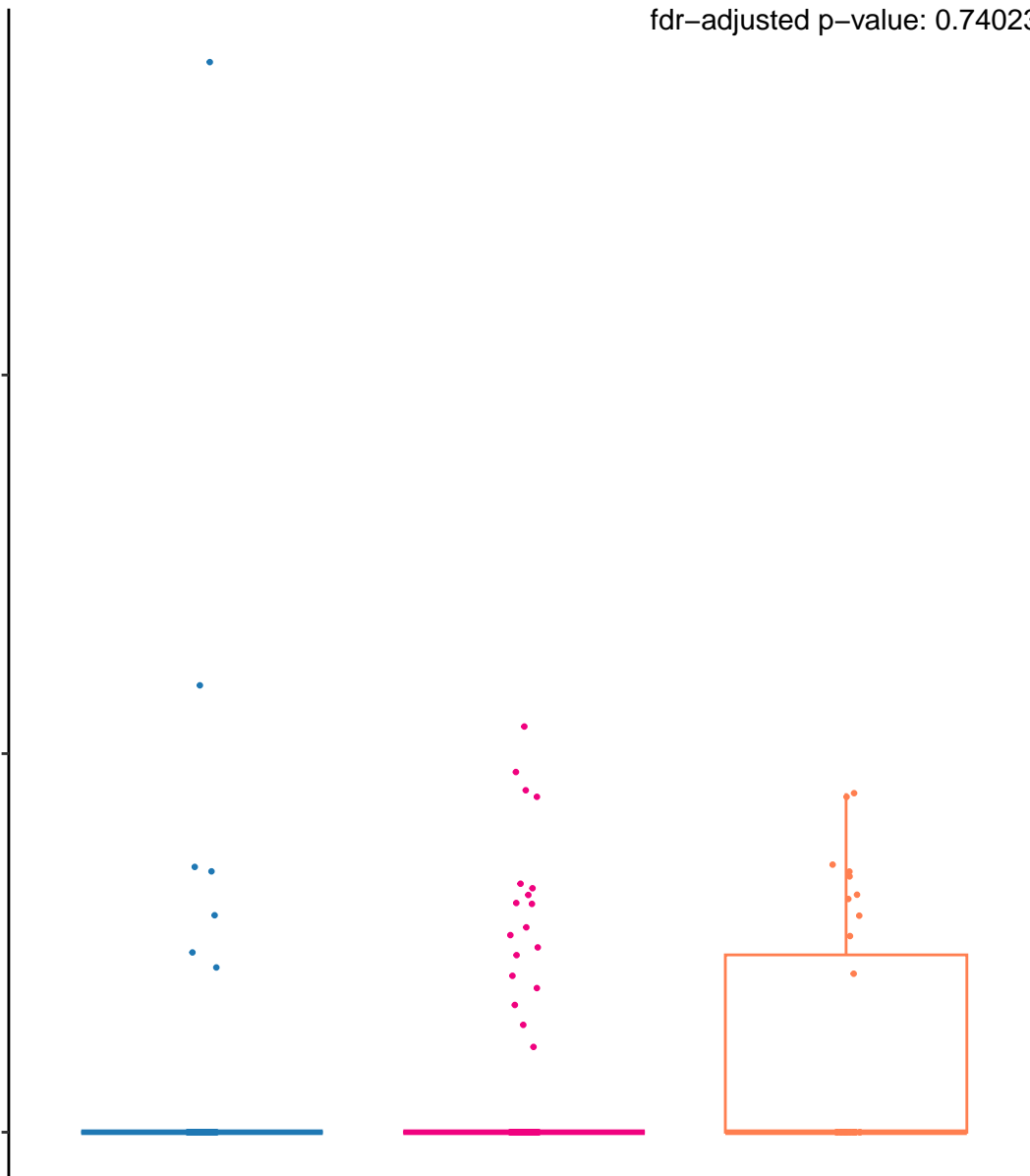
1

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LOCATION

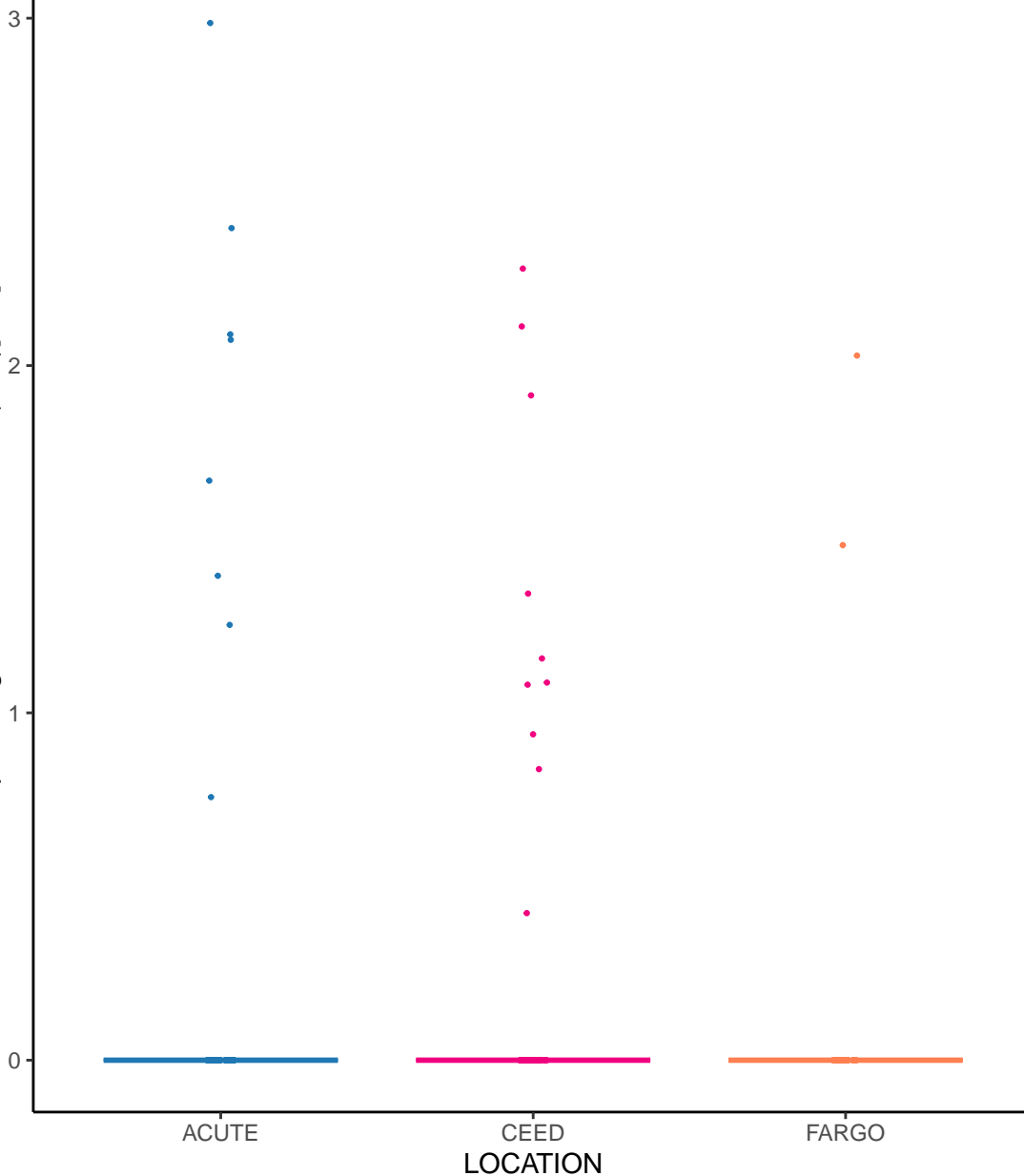


anova p-value: 0.49492

fdr-adjusted p-value: 0.74023

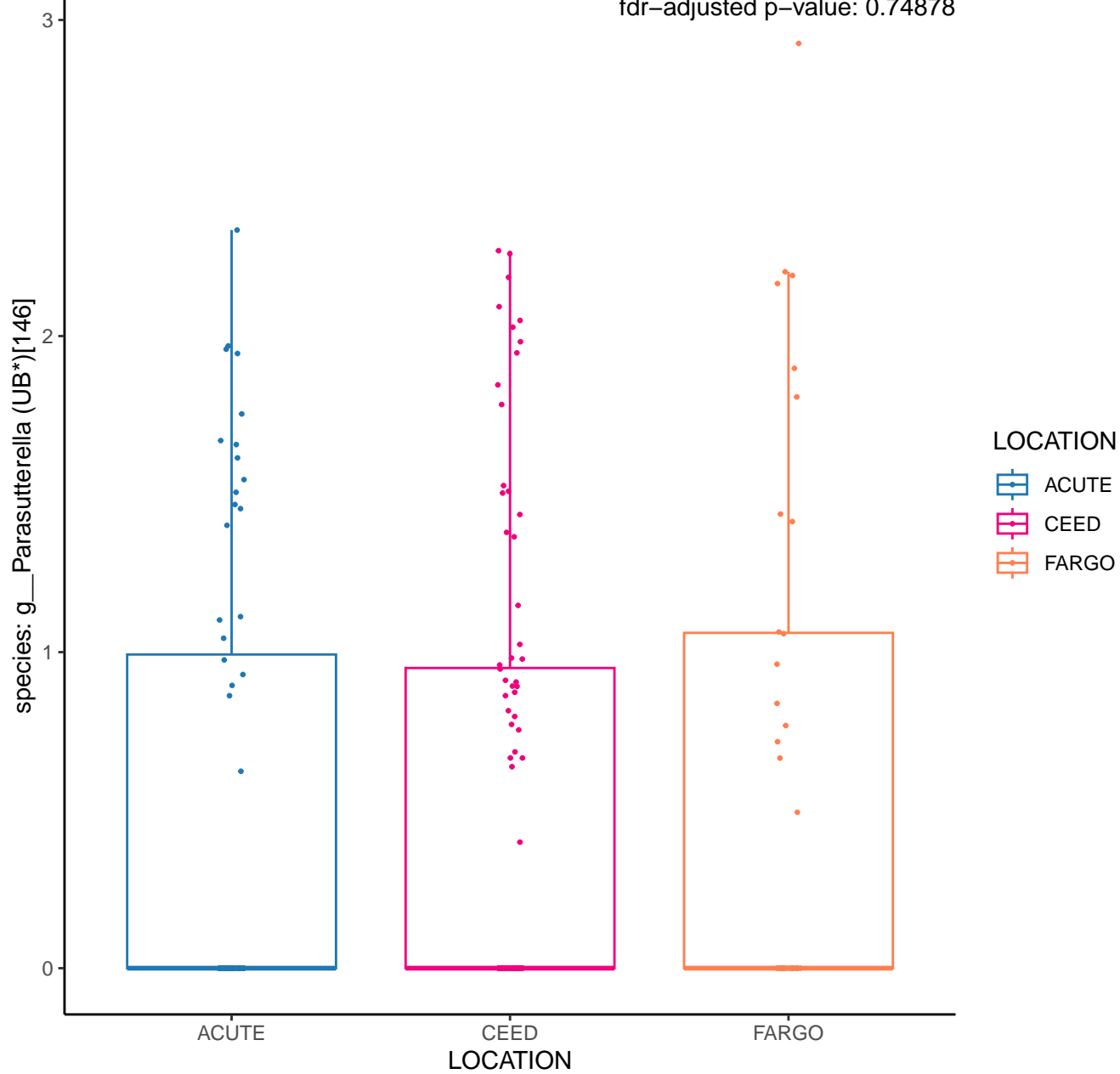
species: g__Acidaminococcus (UB*)[106]

LOCATION



anova p-value: 0.50806

fdr-adjusted p-value: 0.74878



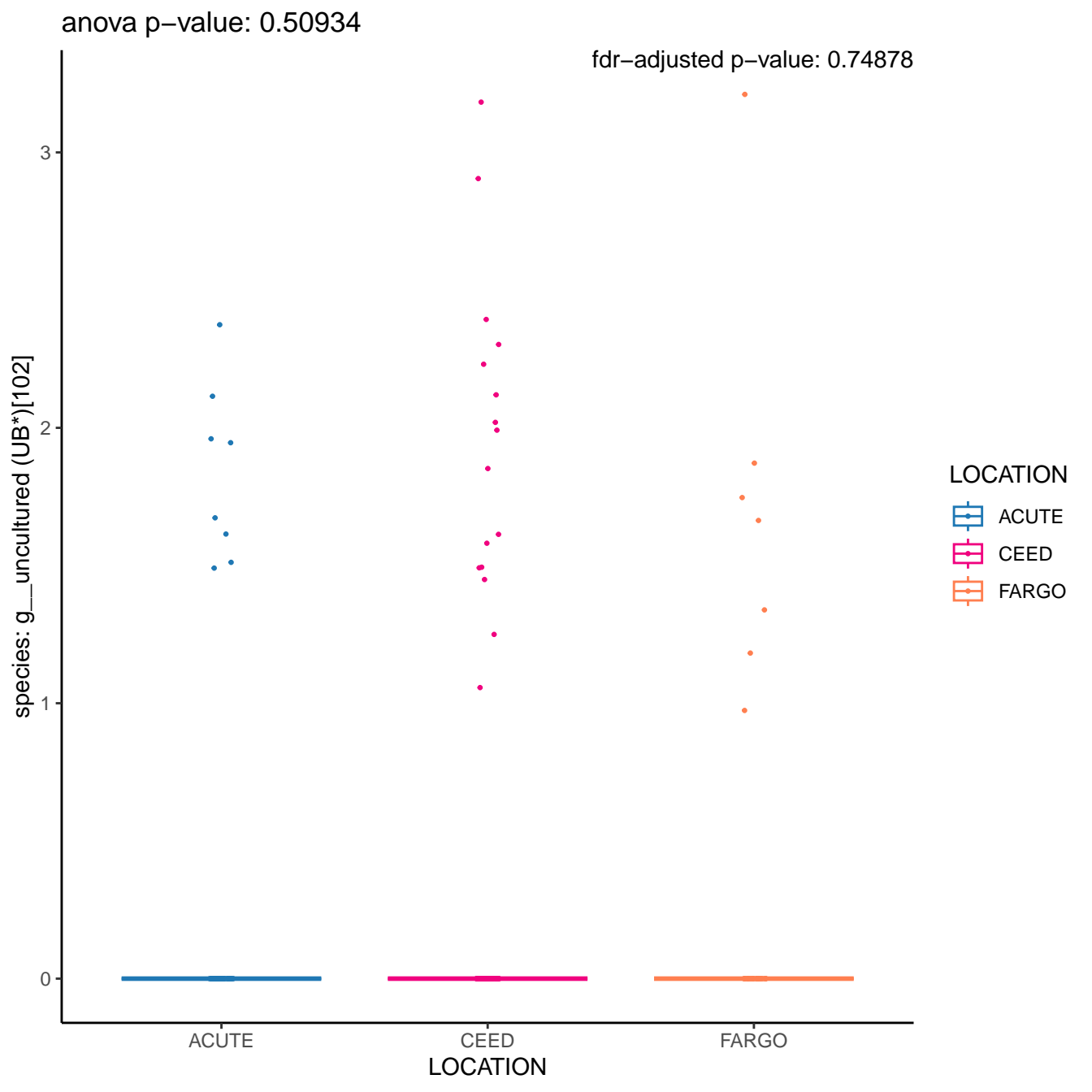
anova p-value: 0.50934

fdr-adjusted p-value: 0.74878

species: g__uncultured (UB*)[102]

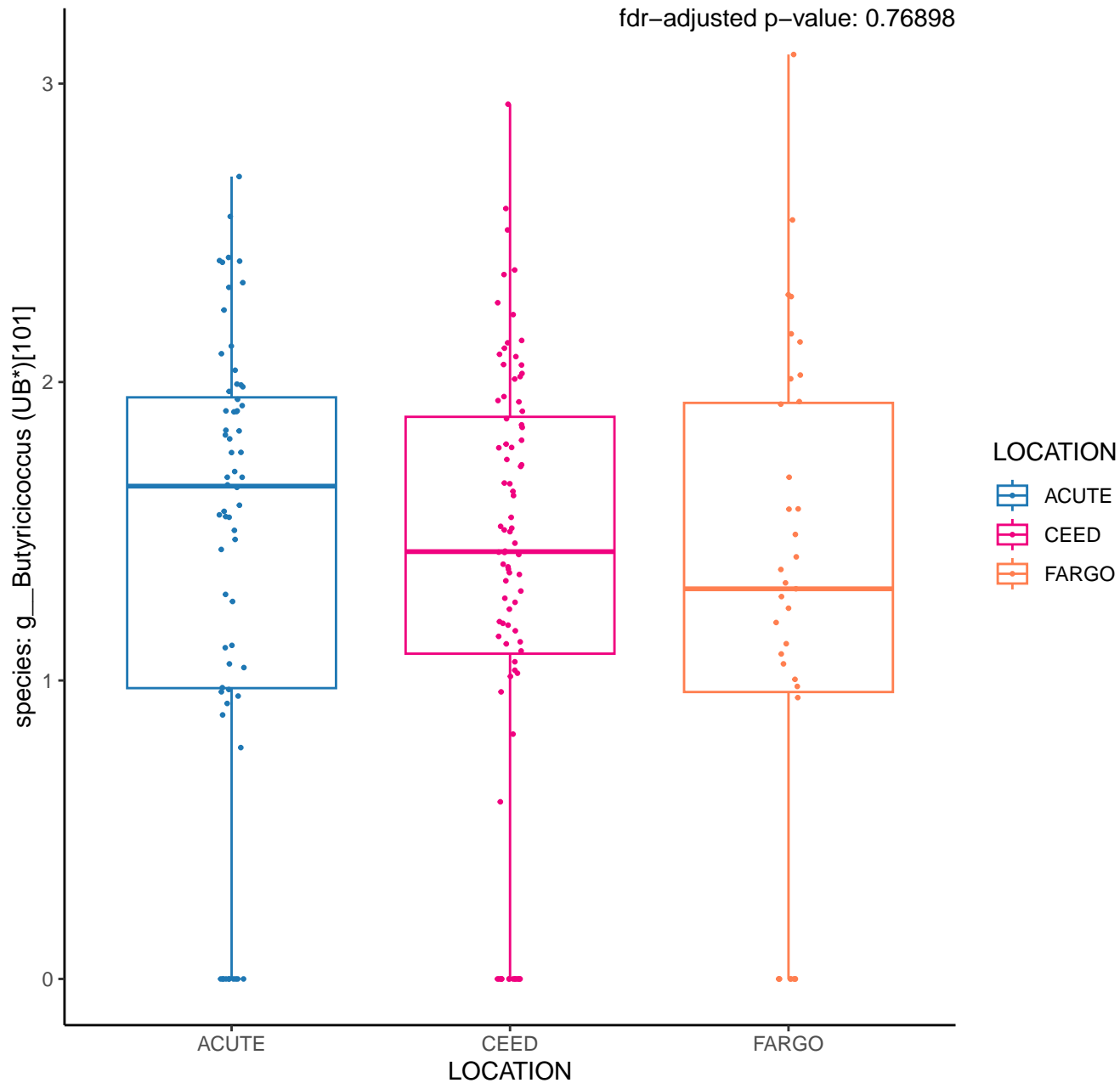
LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.52798

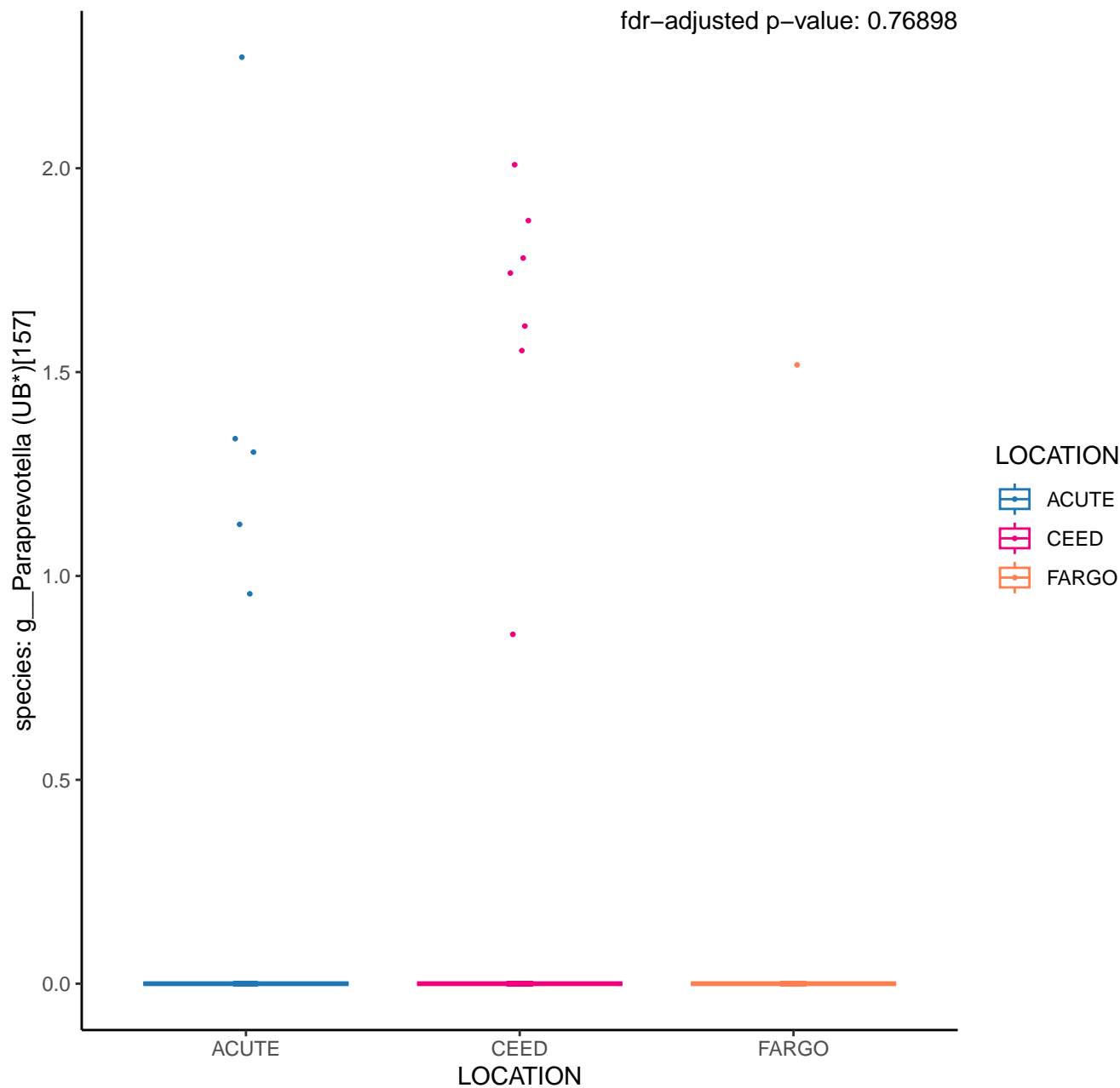
fdr-adjusted p-value: 0.76898



anova p-value: 0.53313

fdr-adjusted p-value: 0.76898

species: g__Paraprevotella (UB*)[157]



anova p-value: 0.55727

fdr-adjusted p-value: 0.76898

species: Roseburia.[17]

LOCATION

ACUTE

CEED

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ACUTE

CEED

FARGO

LOCATION

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anova p-value: 0.55727

fdr-adjusted p-value: 0.76898

species: *Pseudomonas*. [115]

LOCATION

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ACUTE

CEED

FARGO

LOCATION

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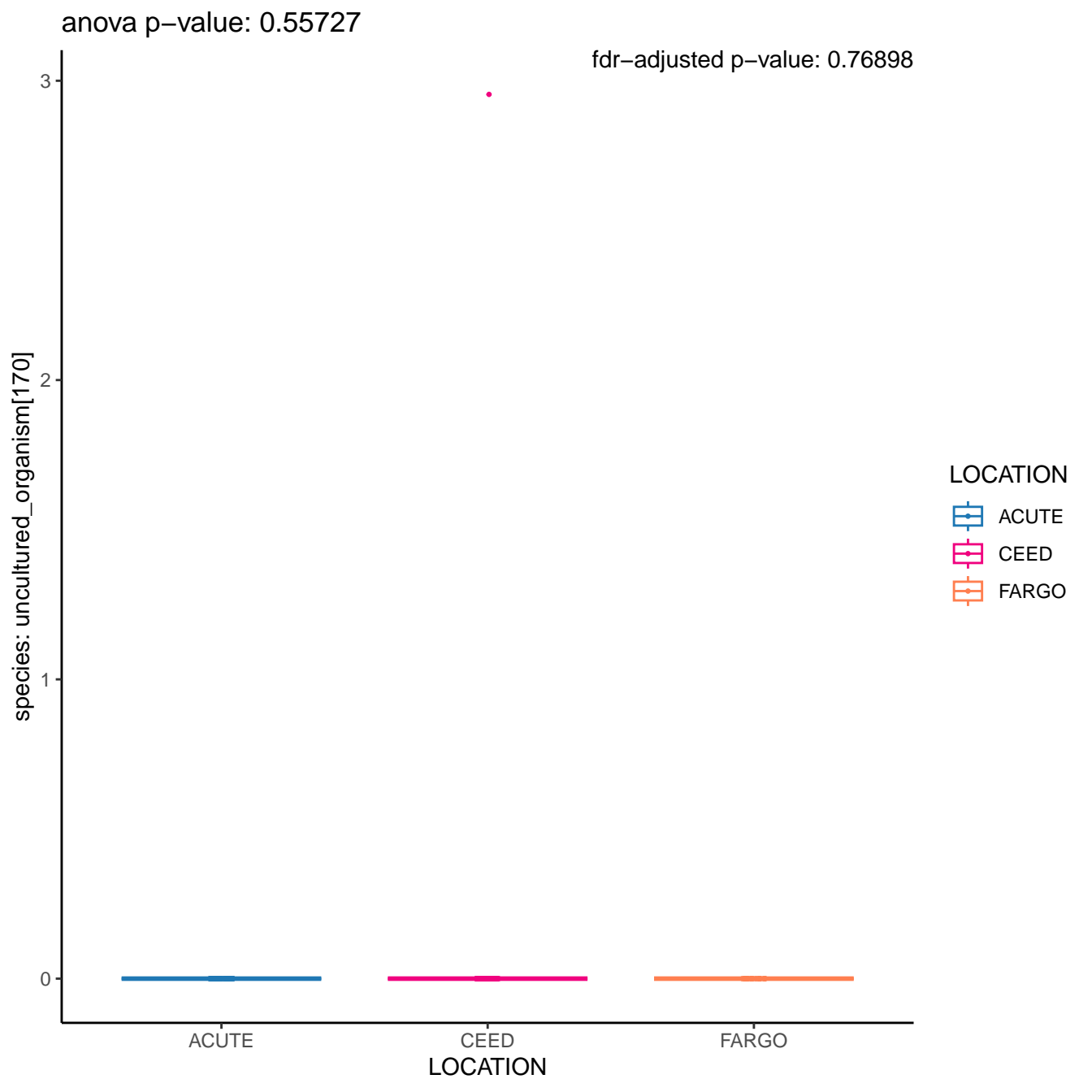
anova p-value: 0.55727

fdr-adjusted p-value: 0.76898

species: uncultured_organism[170]

LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.55727

fdr-adjusted p-value: 0.76898

species: *Klebsiella*.[176]

LOCATION

ACUTE
CEED
FARGO

species: *Klebsiella*.[176]

0

ACUTE

CEED

FARGO

LOCATION

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2

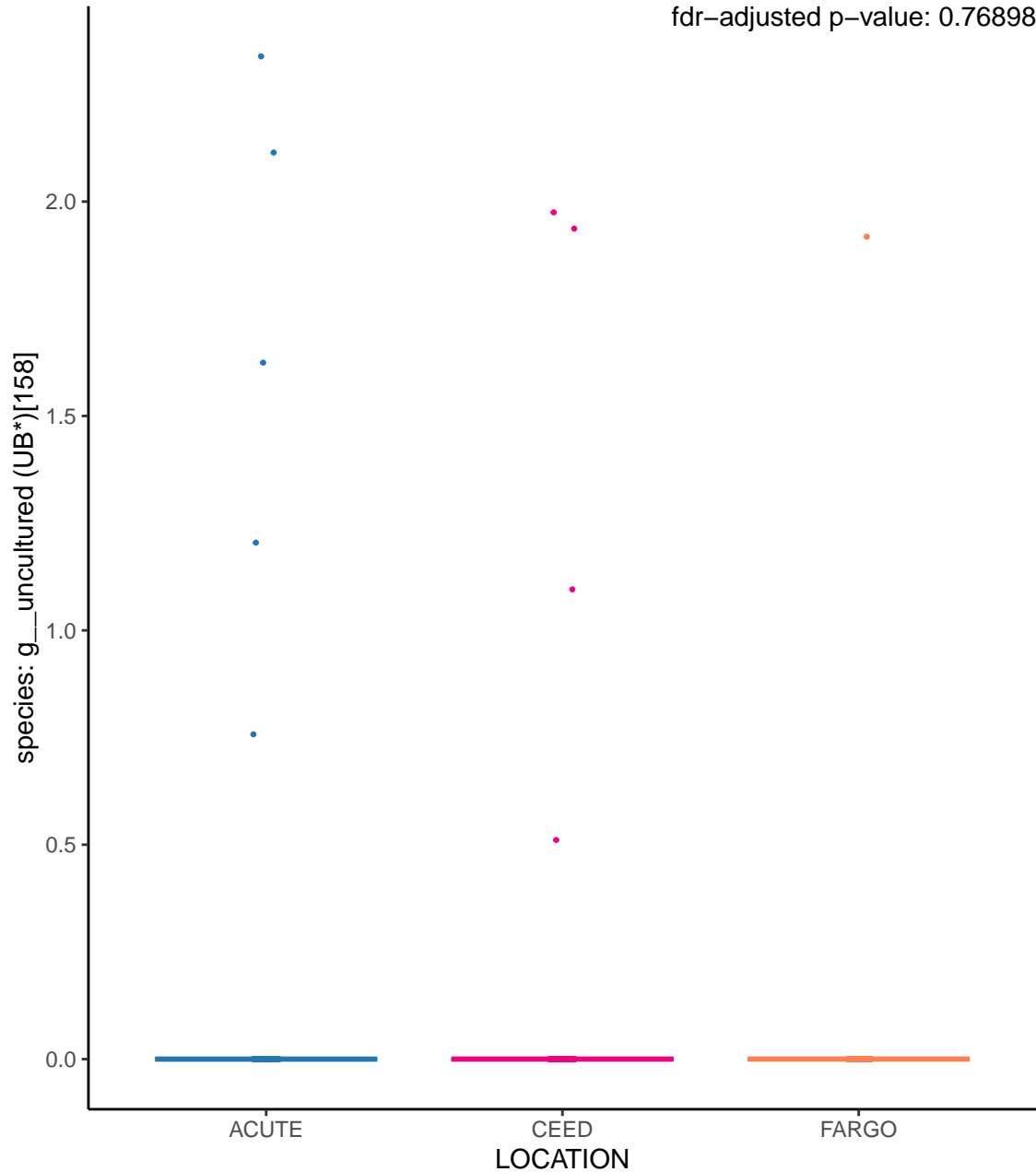
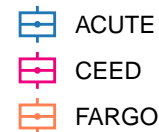
1

anova p-value: 0.55758

fdr-adjusted p-value: 0.76898

species: g__uncultured (UB*)[158]

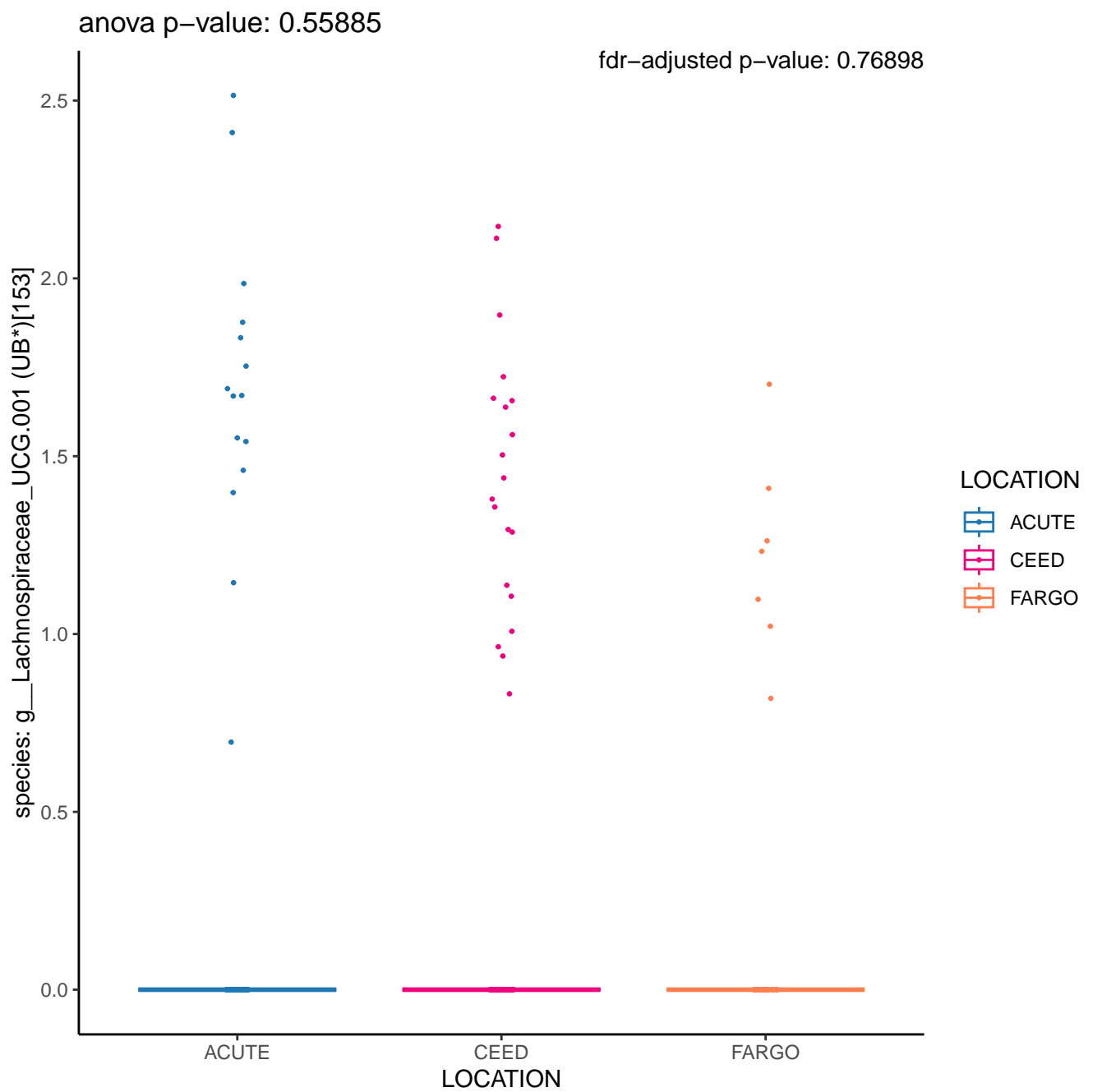
LOCATION



anova p-value: 0.55885

fdr-adjusted p-value: 0.76898

species: g__Lachnospiraceae_UCG.001 (UB*)[153]



anova p-value: 0.58306

fdr-adjusted p-value: 0.79149

species: *Eggerthella*. [83]

LOCATION

ACUTE

CEED

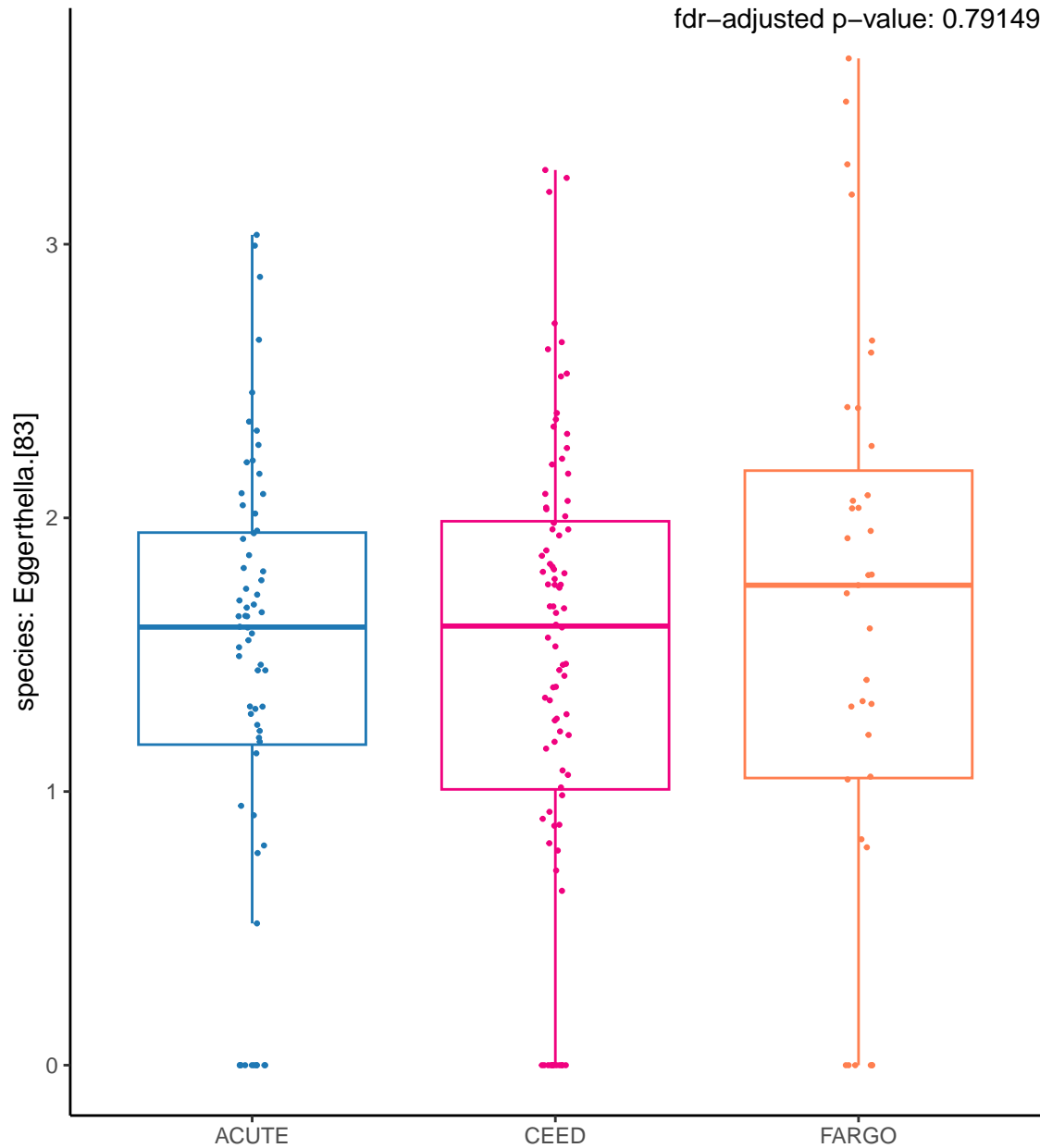
FARGO

ACUTE

CEED

FARGO

LOCATION



fdr-adjusted p-value: 0.79149

anova p-value: 0.61553

fdr-adjusted p-value: 0.82711

species: Cloacibacillus.[173]

LOCATION

ACUTE
CEED
FARGO

2.0
1.5
1.0
0.5
0.0

ACUTE

CEED

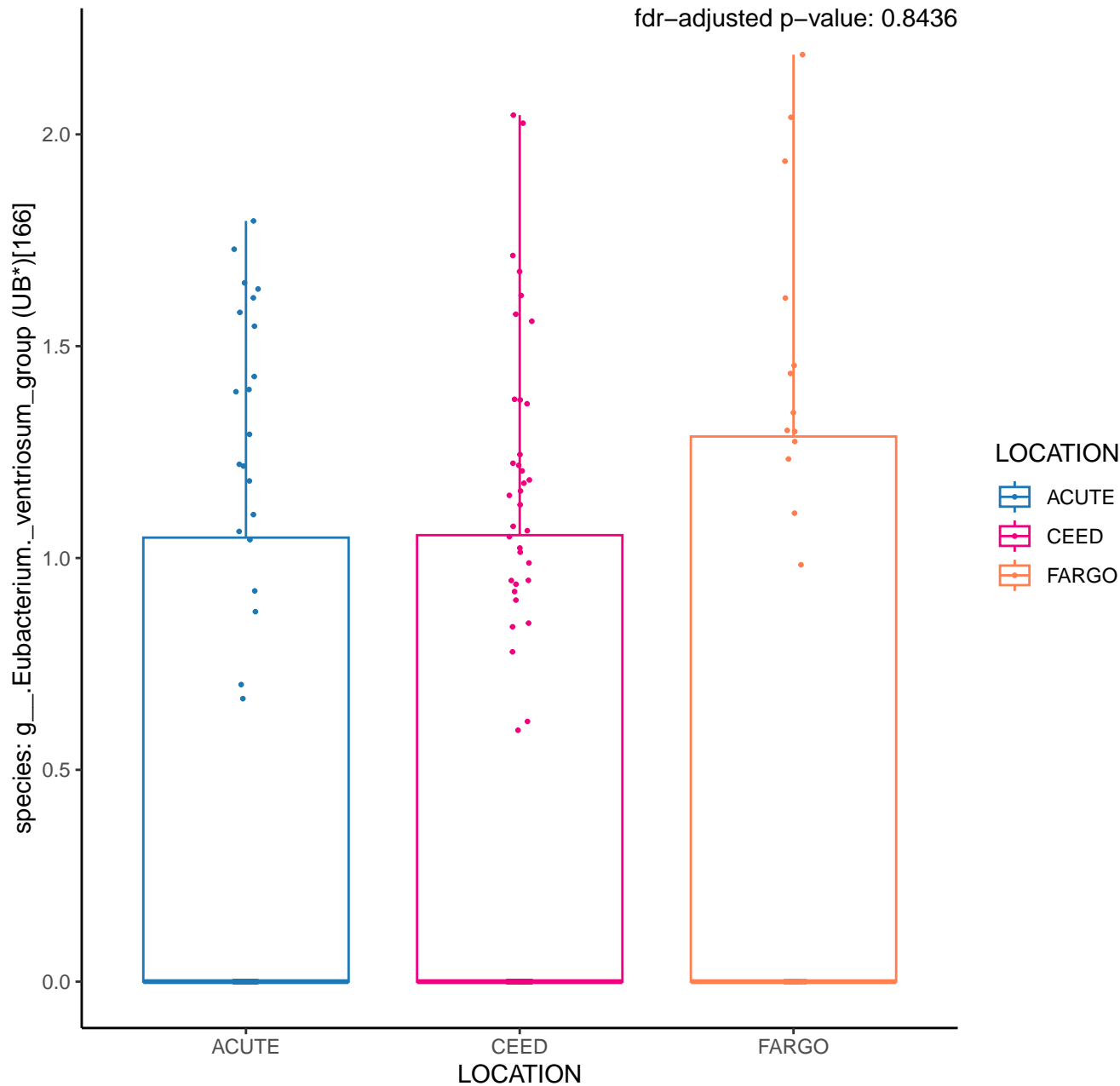
FARGO

LOCATION



anova p-value: 0.63525

fdr-adjusted p-value: 0.8436



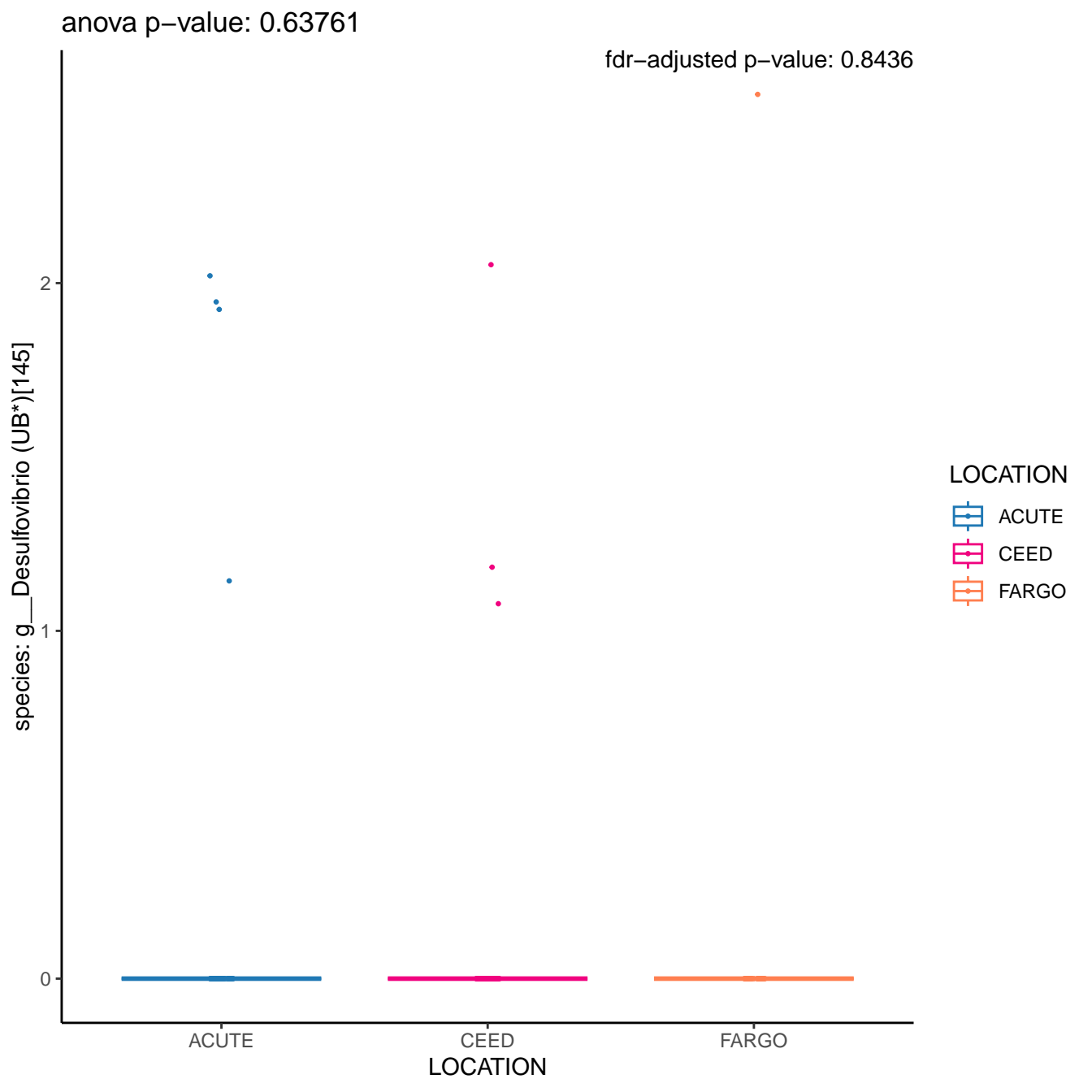
anova p-value: 0.63761

fdr-adjusted p-value: 0.8436

species: g__Desulfovibrio (UB*)[145]

LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.65486

fdr-adjusted p-value: 0.84395

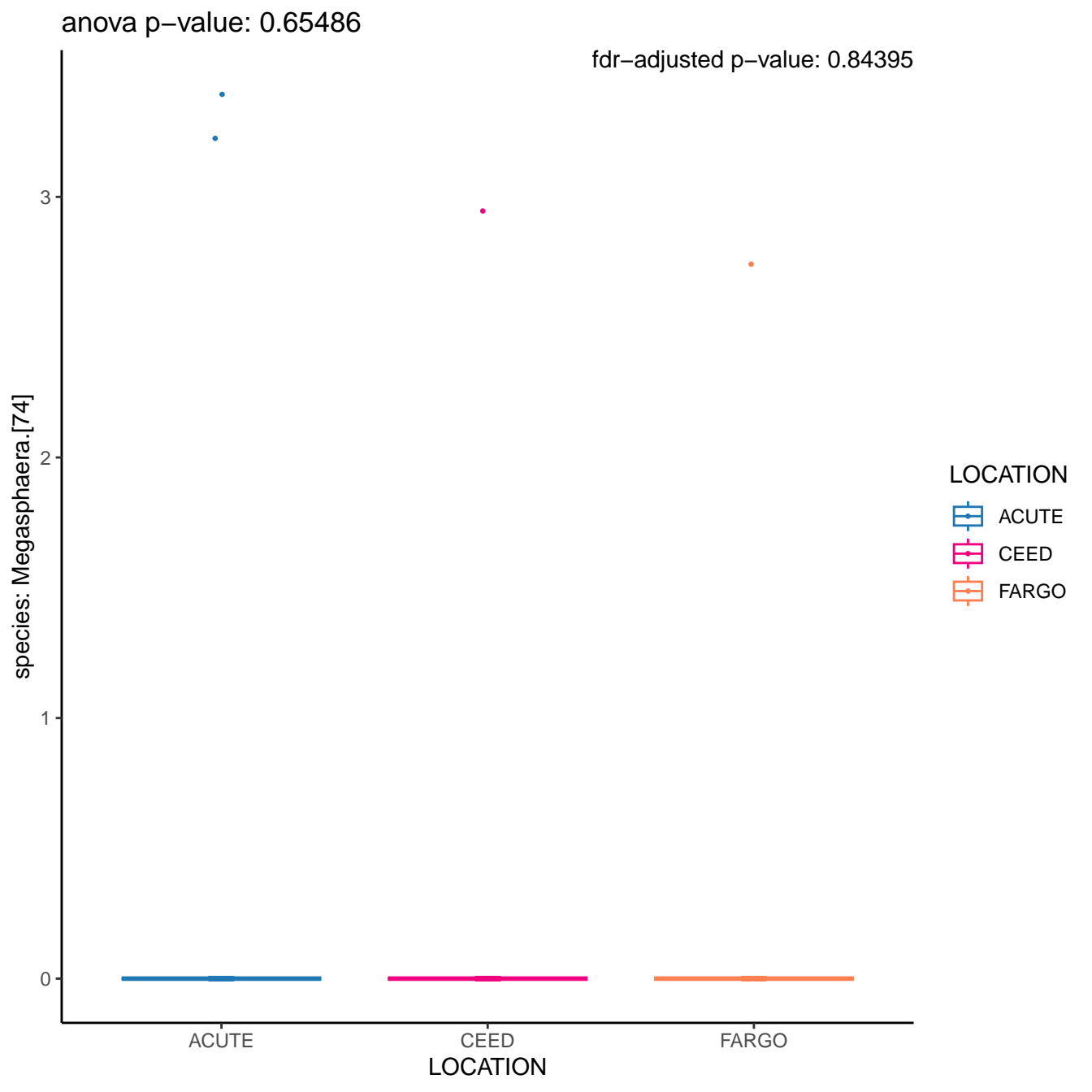
species: Megasphaera.[74]

LOCATION

ACUTE

CEED

FARGO



anova p-value: 0.65752

fdr-adjusted p-value: 0.84395

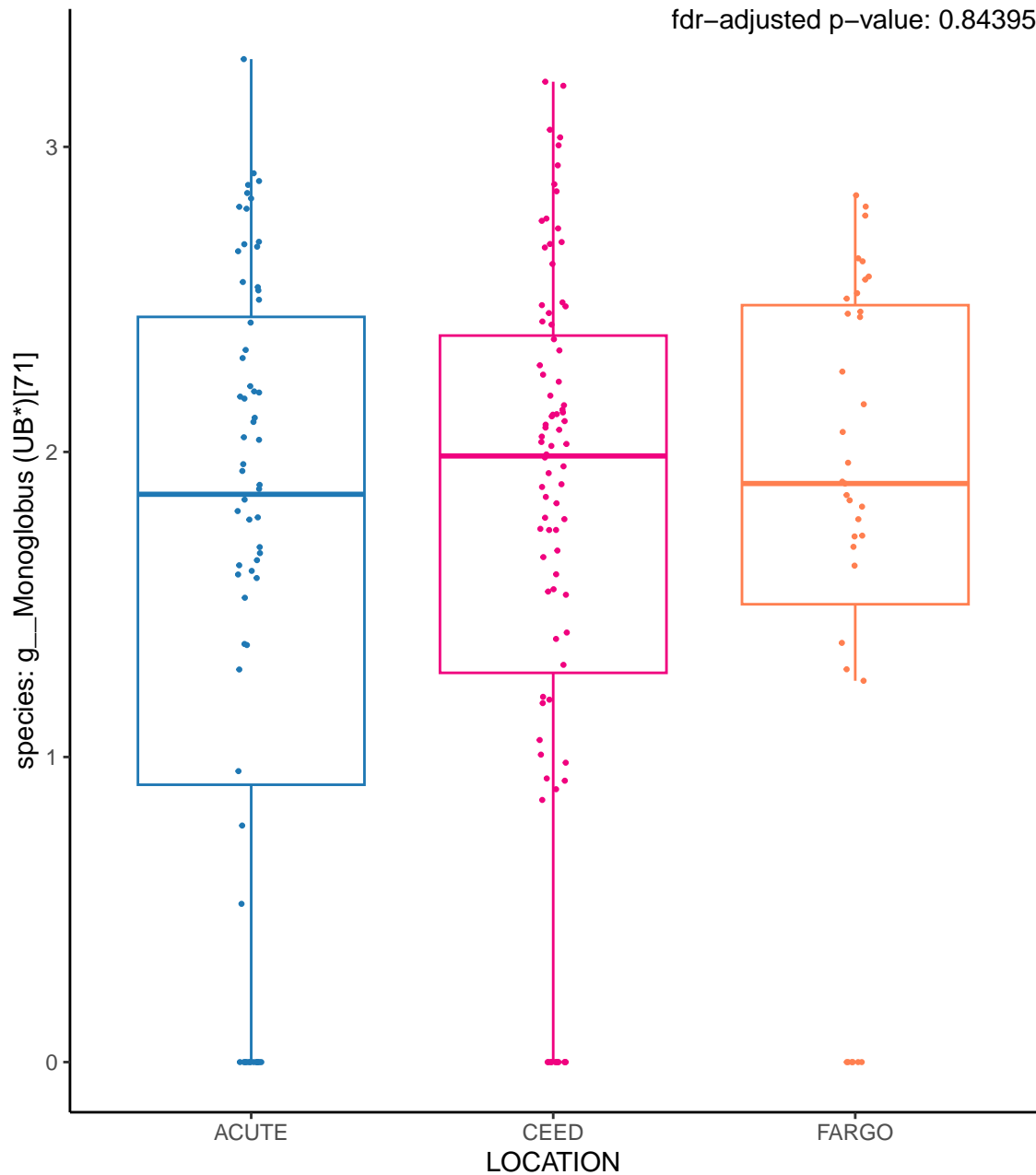
species: g__Monoglobus (UB*)[71]

LOCATION

ACUTE

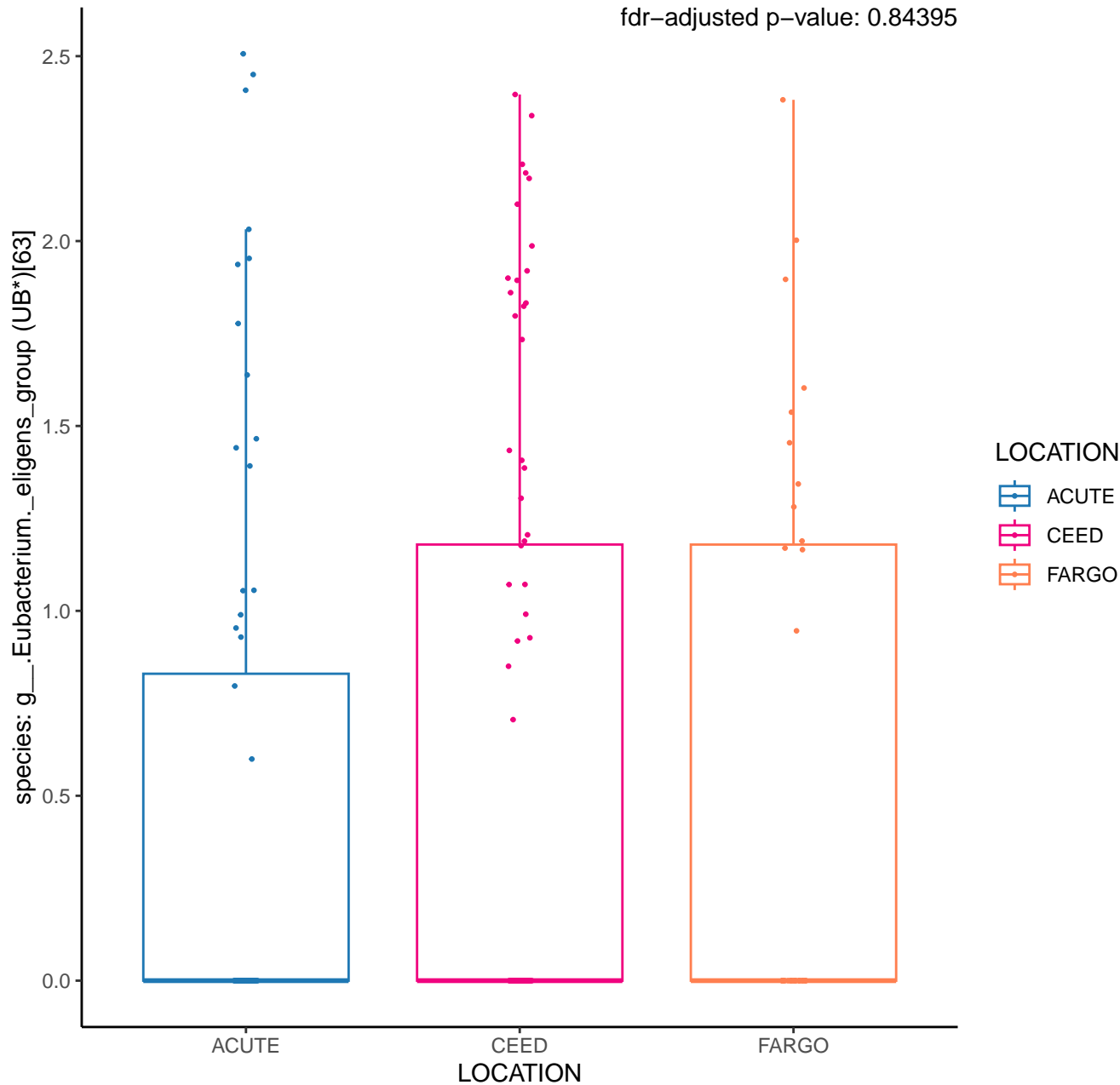
CEED

FARGO

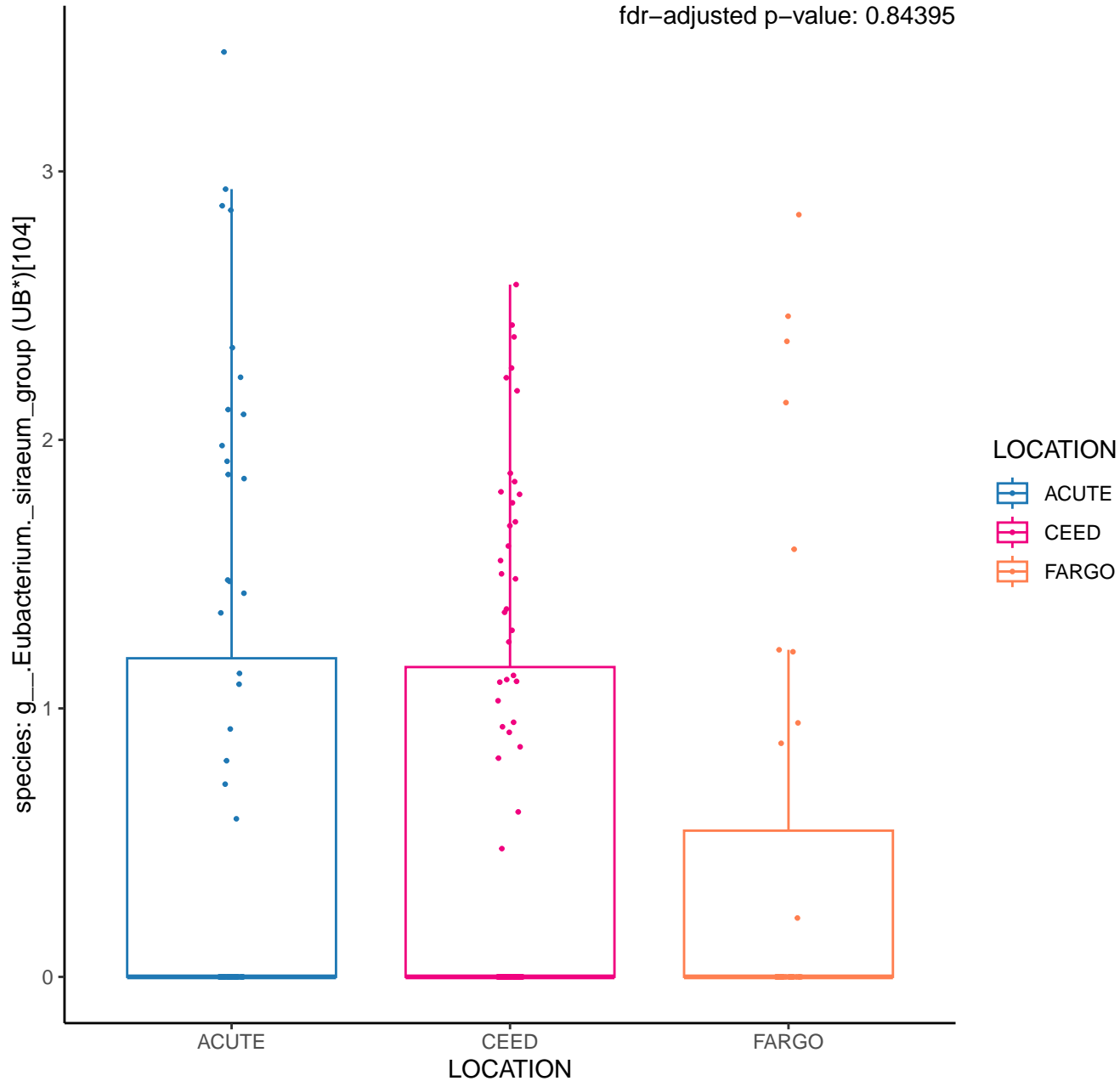


anova p-value: 0.6596

fdr-adjusted p-value: 0.84395



fdr-adjusted p-value: 0.84395



anova p-value: 0.66241

fdr-adjusted p-value: 0.84395

species: .Eubacterium._eligans_group.[136]

LOCATION

ACUTE

CEED

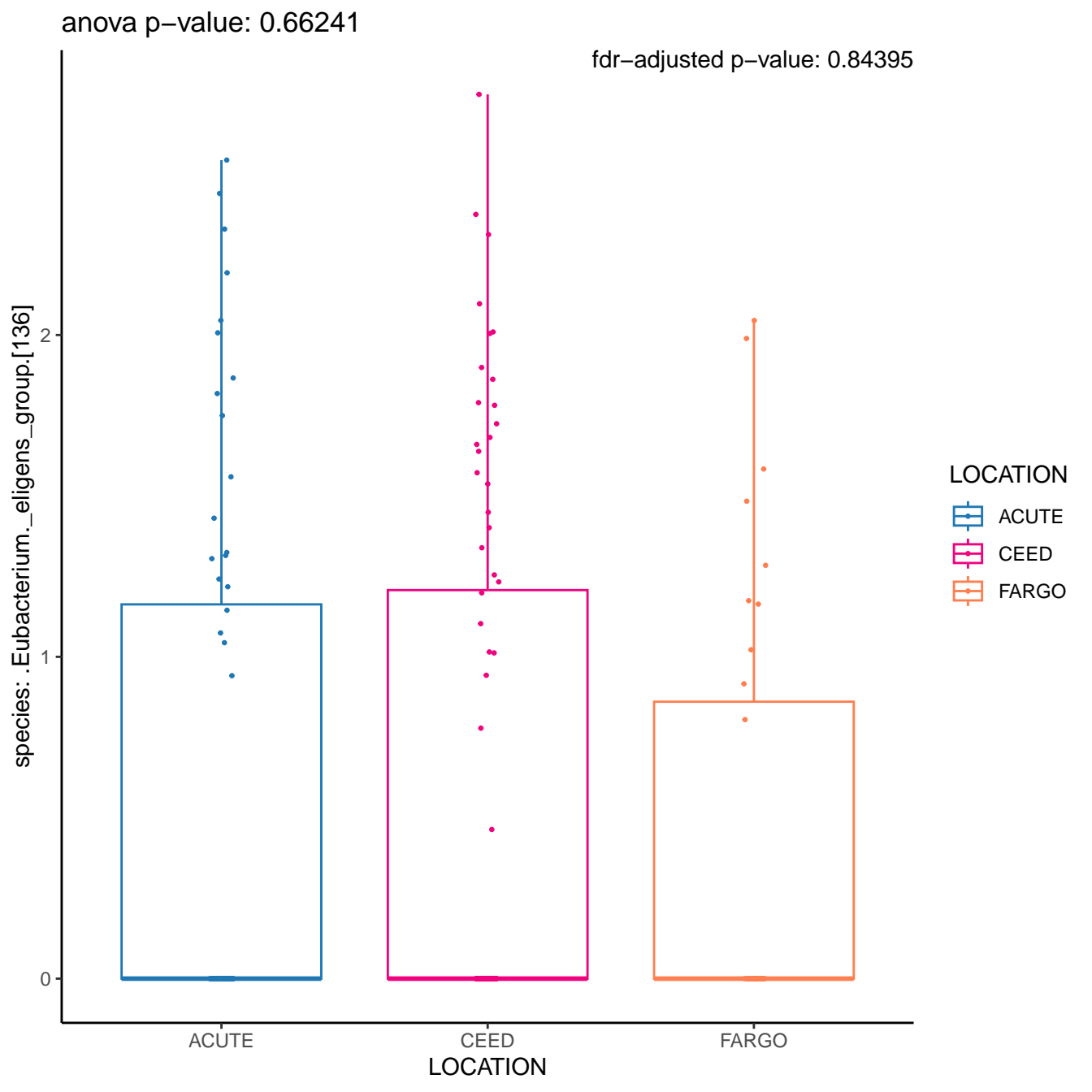
FARGO

ACUTE

CEED

FARGO

LOCATION



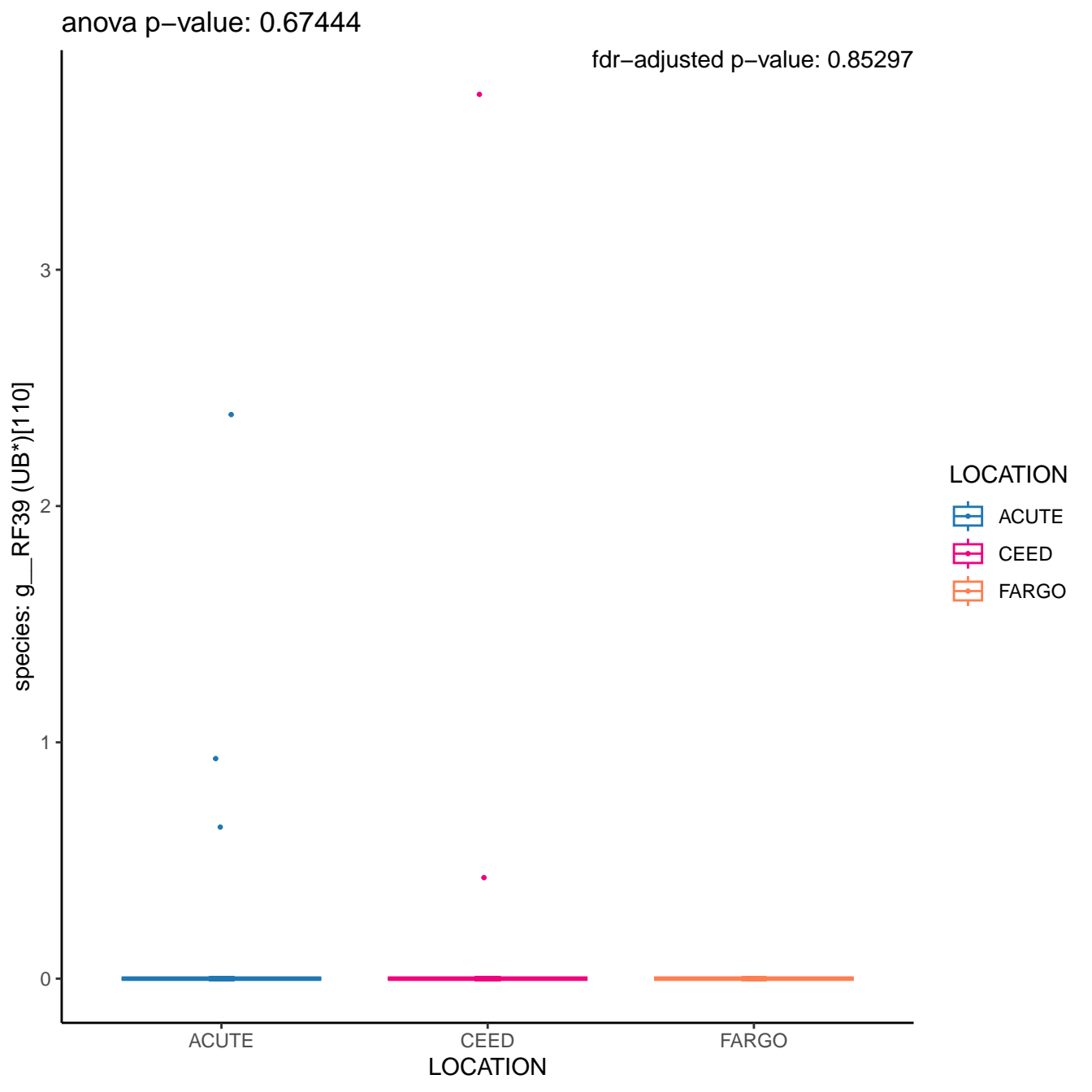
anova p-value: 0.67444

fdr-adjusted p-value: 0.85297

species: g__RF39 (UB*)[110]

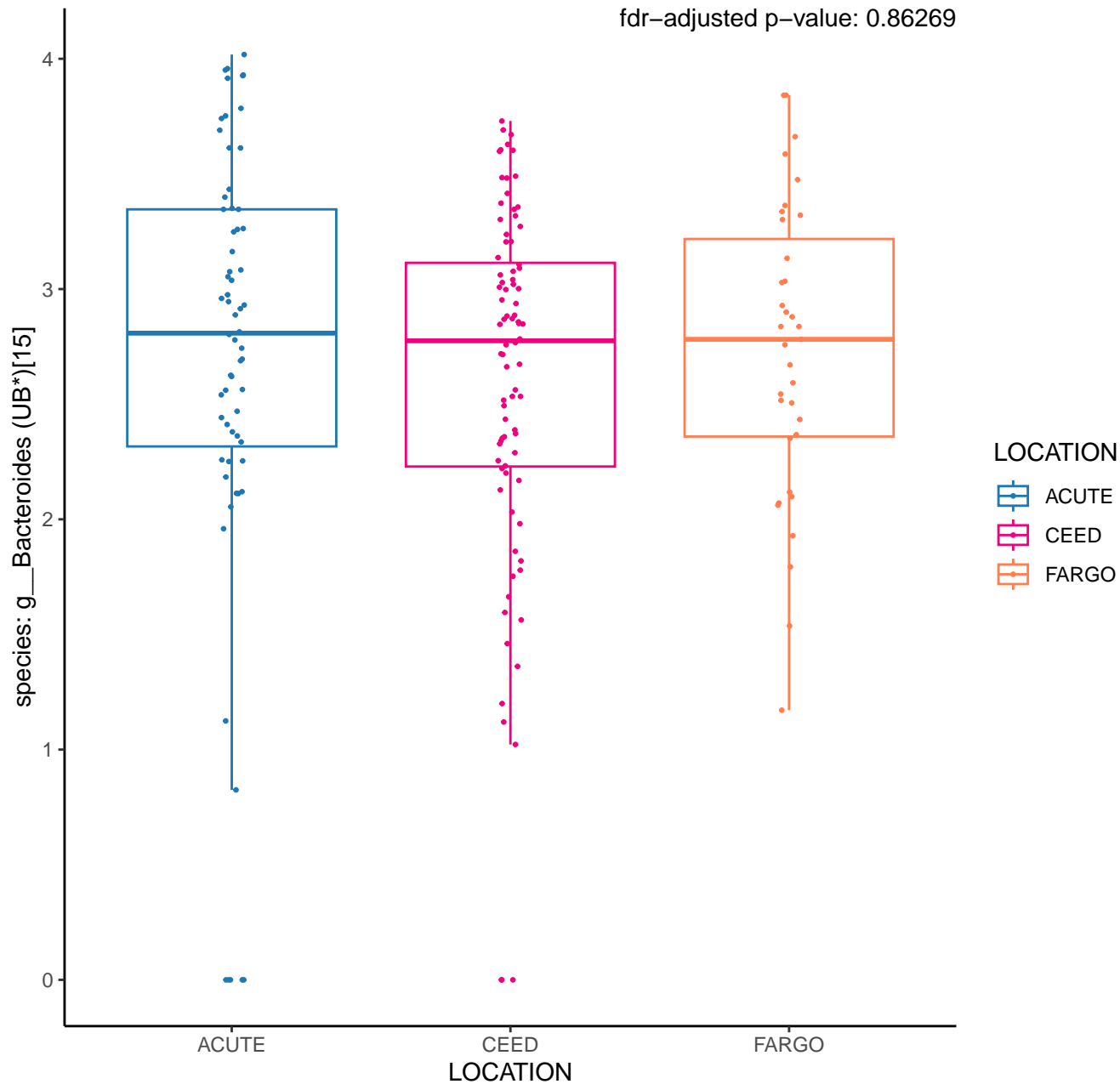
LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.68714

fdr-adjusted p-value: 0.86269



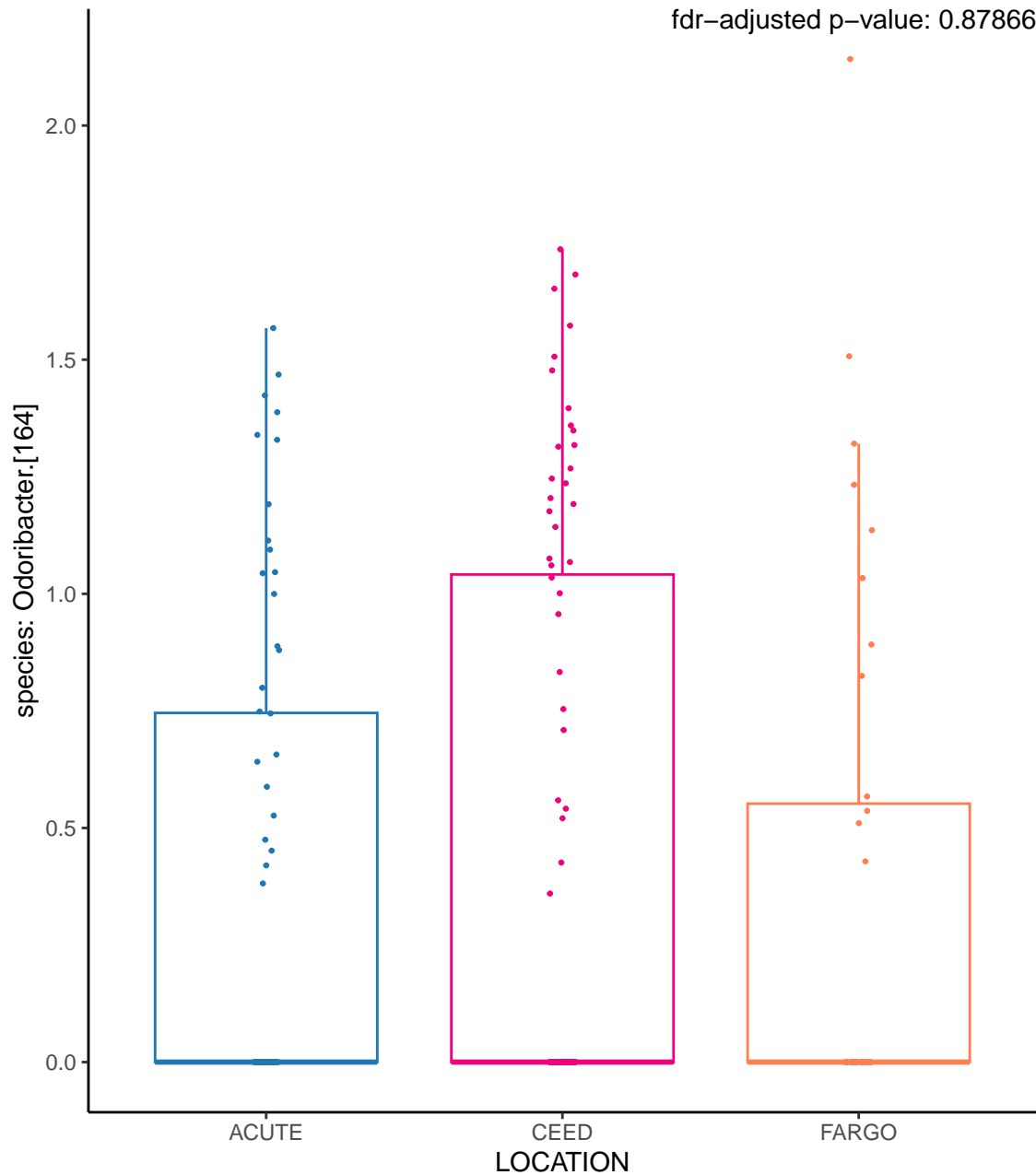
anova p-value: 0.70497

fdr-adjusted p-value: 0.87866

species: Odoribacter.[164]

LOCATION

ACUTE
CEED
FARGO



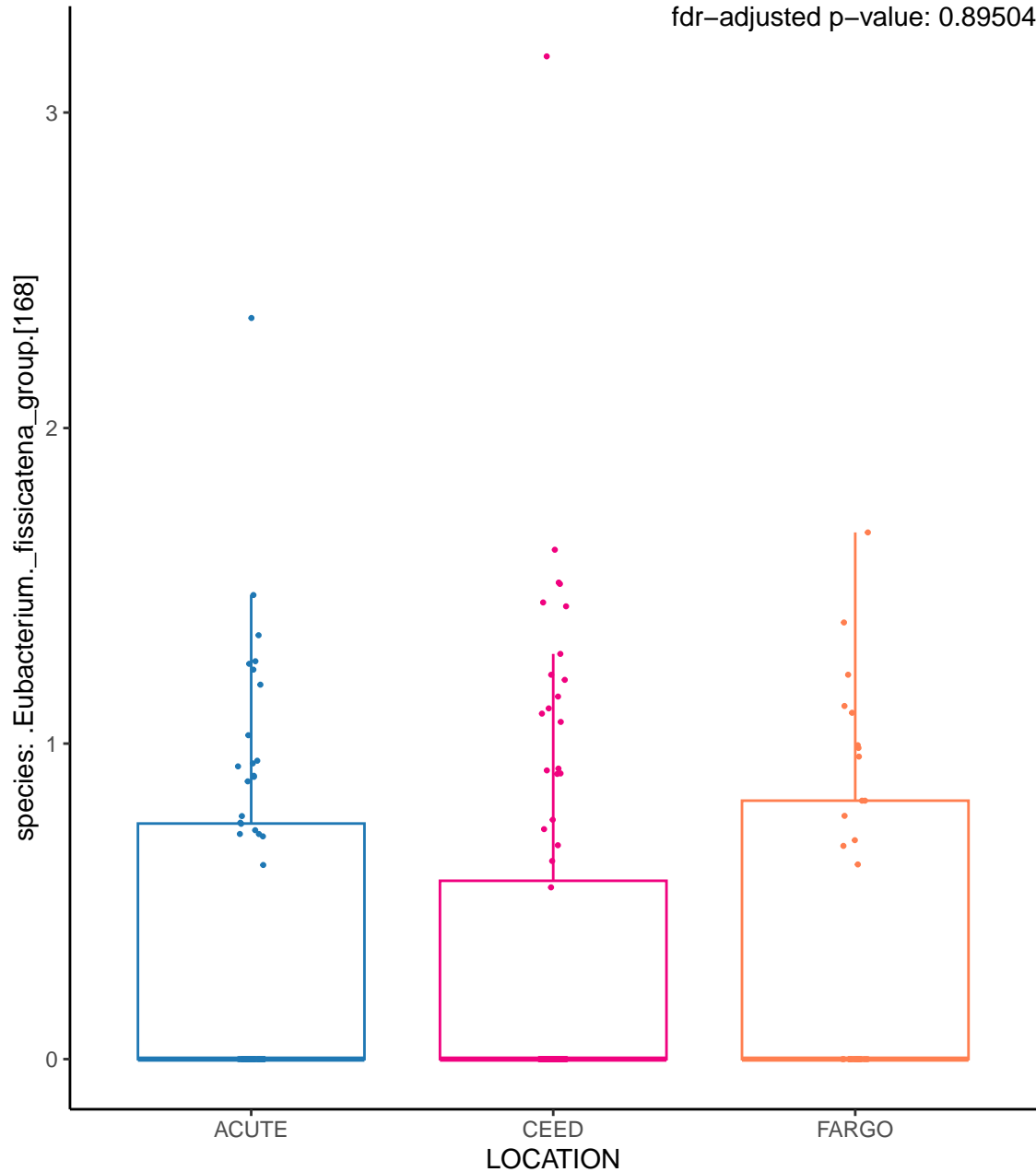
anova p-value: 0.72332

fdr-adjusted p-value: 0.89504

species: *Eubacterium._fissicatena_group.[168]*

LOCATION

ACUTE
CEED
FARGO



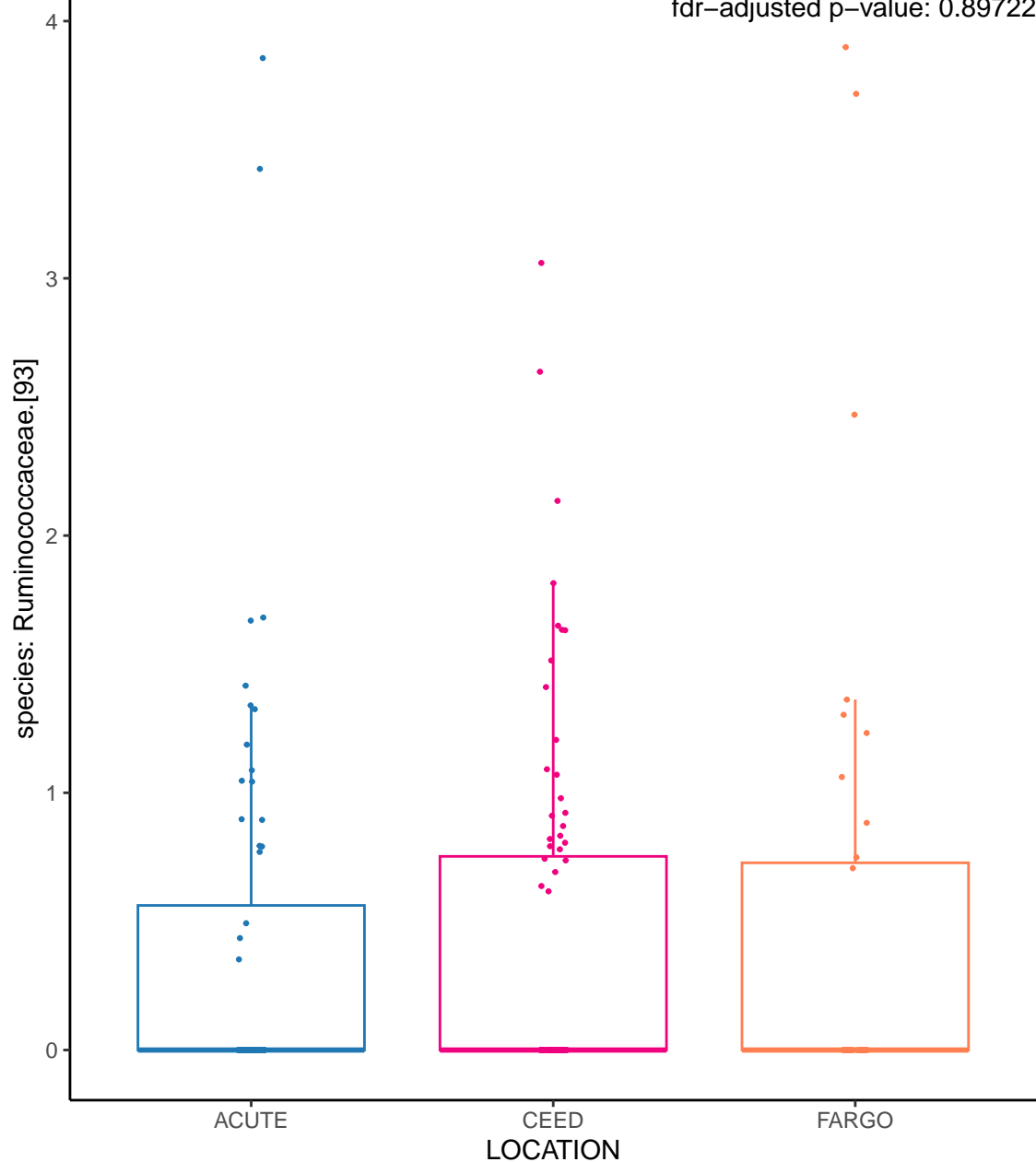
anova p-value: 0.73429

fdr-adjusted p-value: 0.89722

species: Ruminococcaceae.[93]

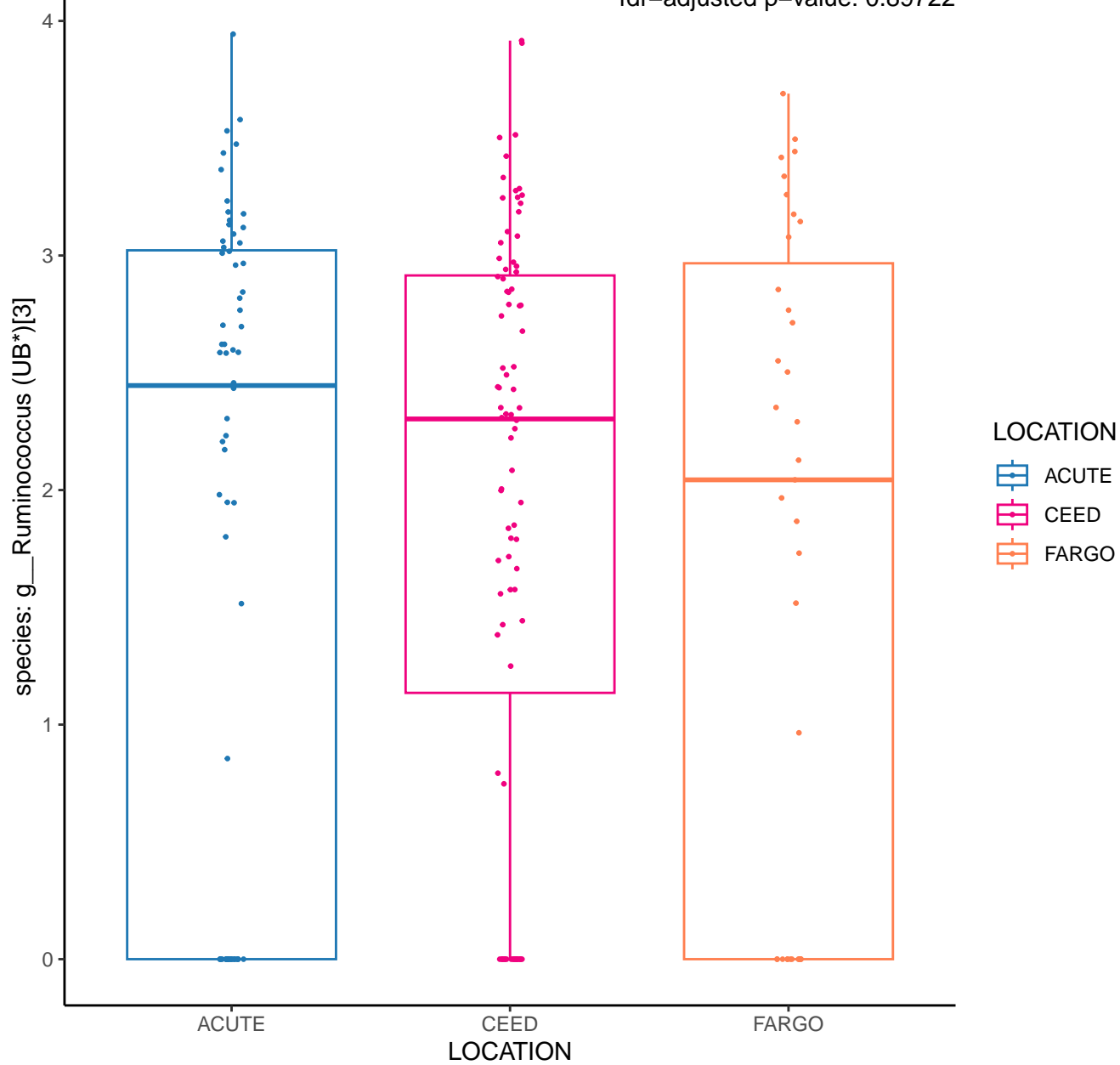
LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.73551

fdr-adjusted p-value: 0.89722



anova p-value: 0.74895

fdr-adjusted p-value: 0.90718

species: g__Incertae_Sedis (UB*)[76]

LOCATION

ACUTE

CEED

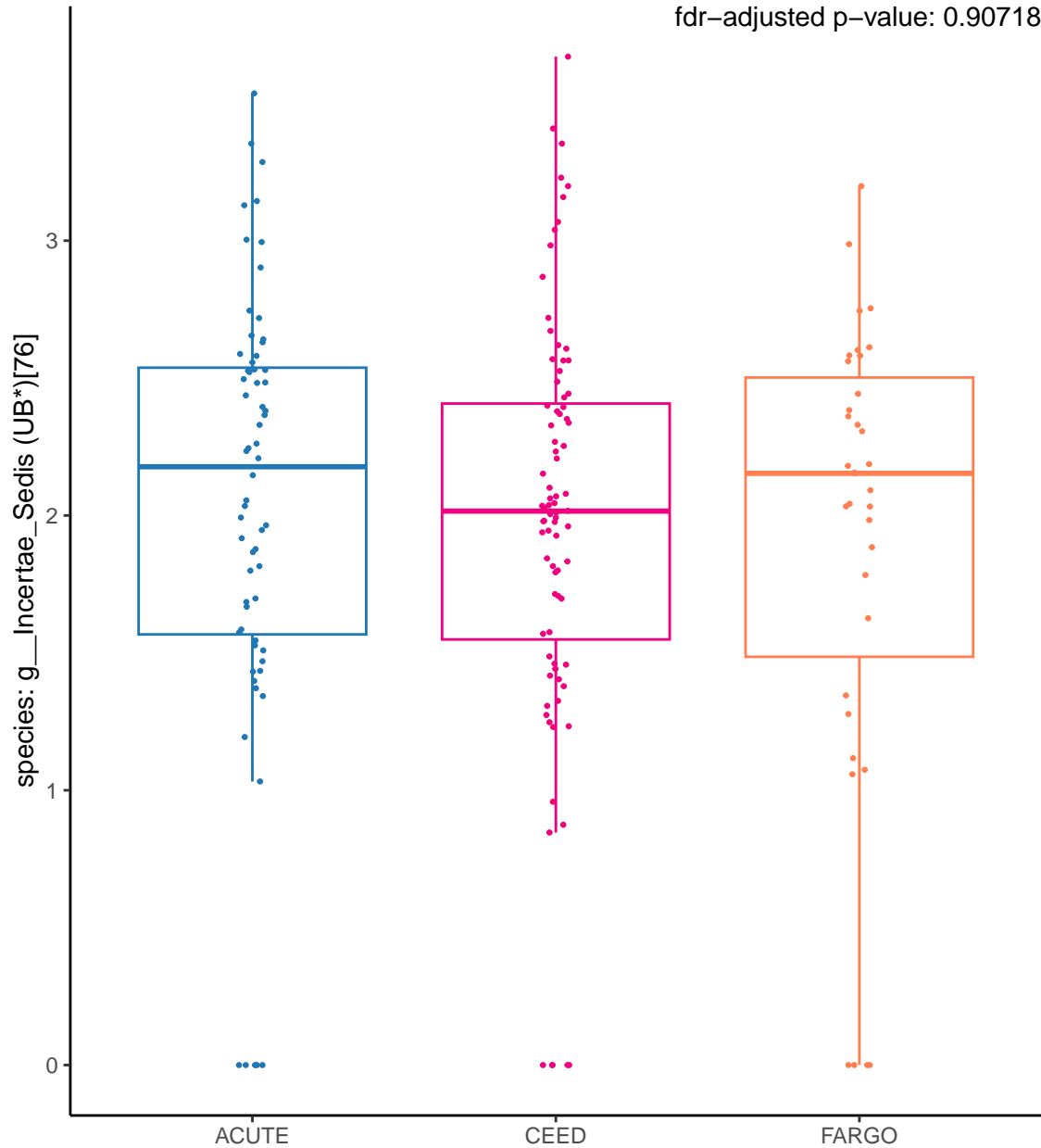
FARGO

ACUTE

CEED

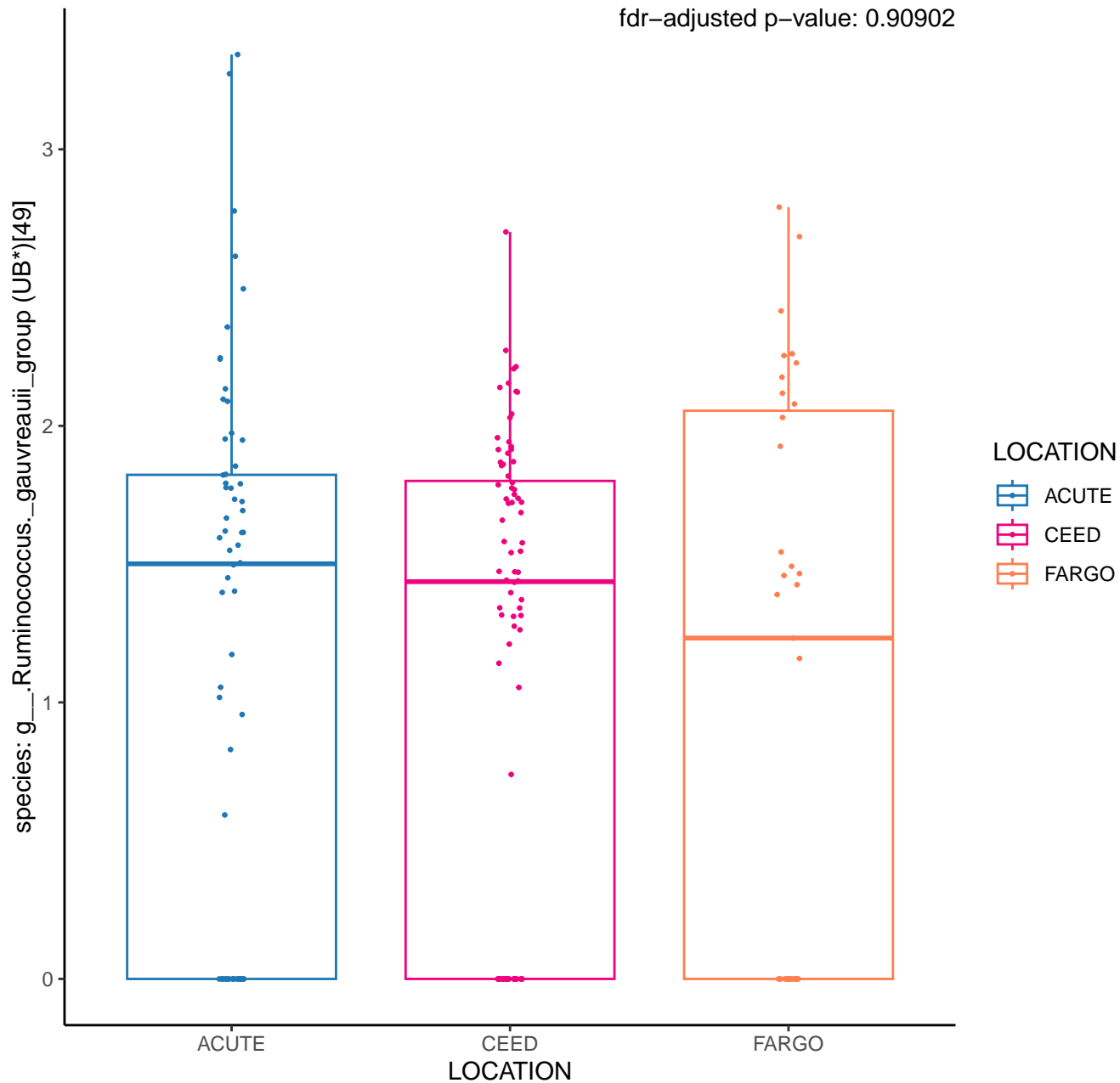
FARGO

LOCATION



anova p-value: 0.75632

fdr-adjusted p-value: 0.90902



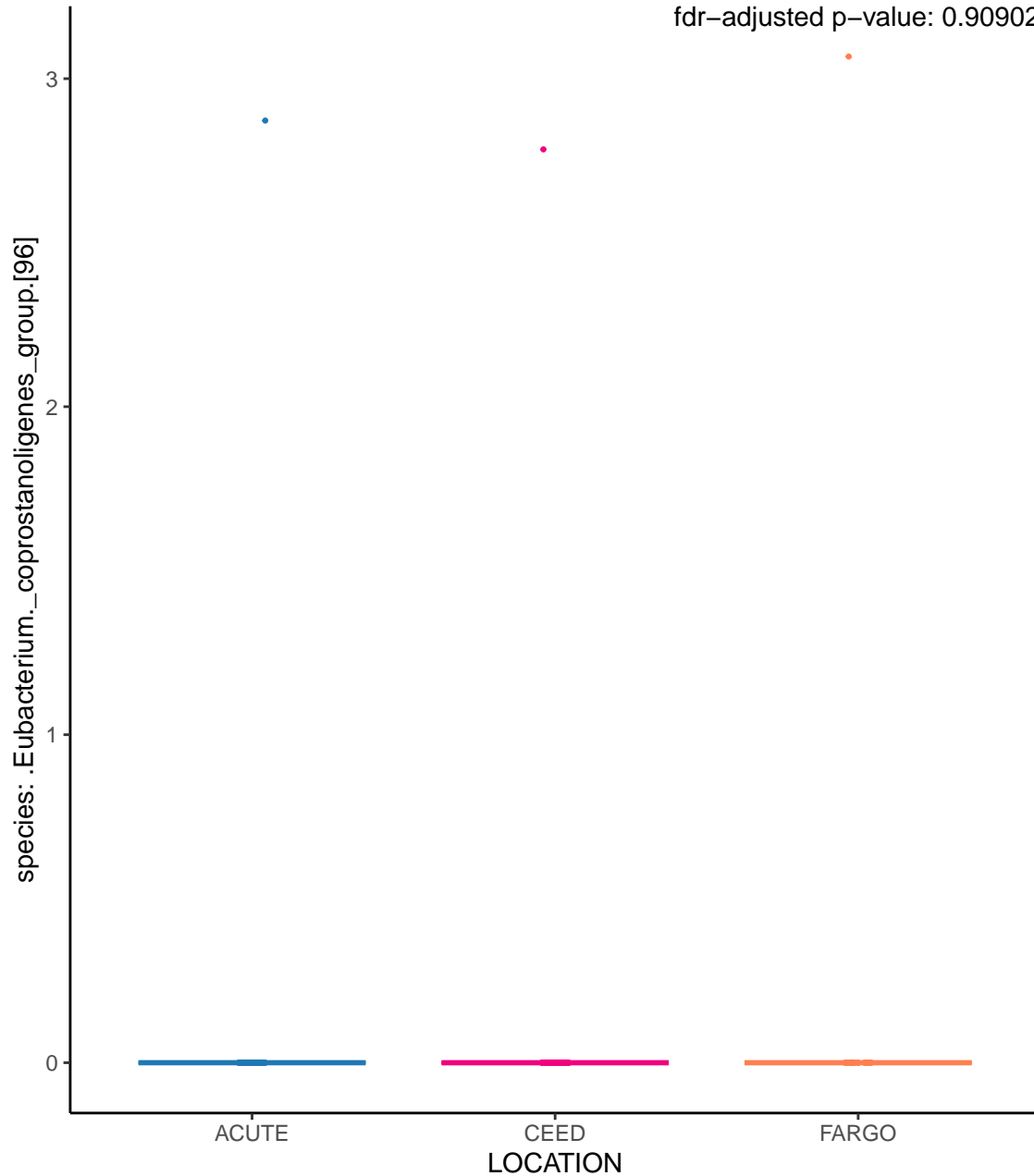
anova p-value: 0.76544

fdr-adjusted p-value: 0.90902

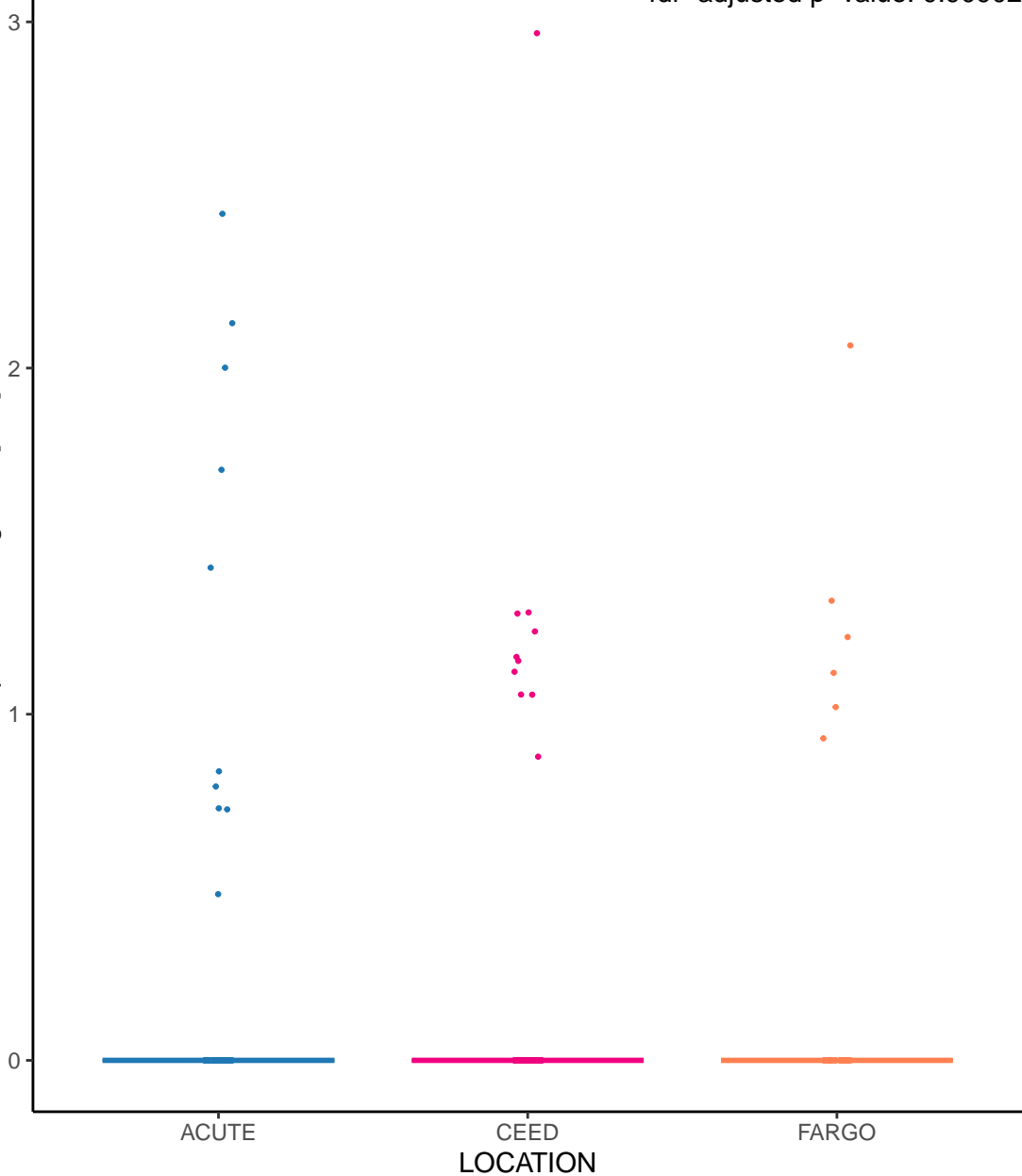
species: .Eubacterium._coprostanoligenes_group.[96]

LOCATION

ACUTE
CEED
FARGO



fdr-adjusted p-value: 0.90902



anova p-value: 0.77444

fdr-adjusted p-value: 0.90902

species: Agathobacter.[54]

LOCATION

ACUTE

CEED

FARGO

0

1

2

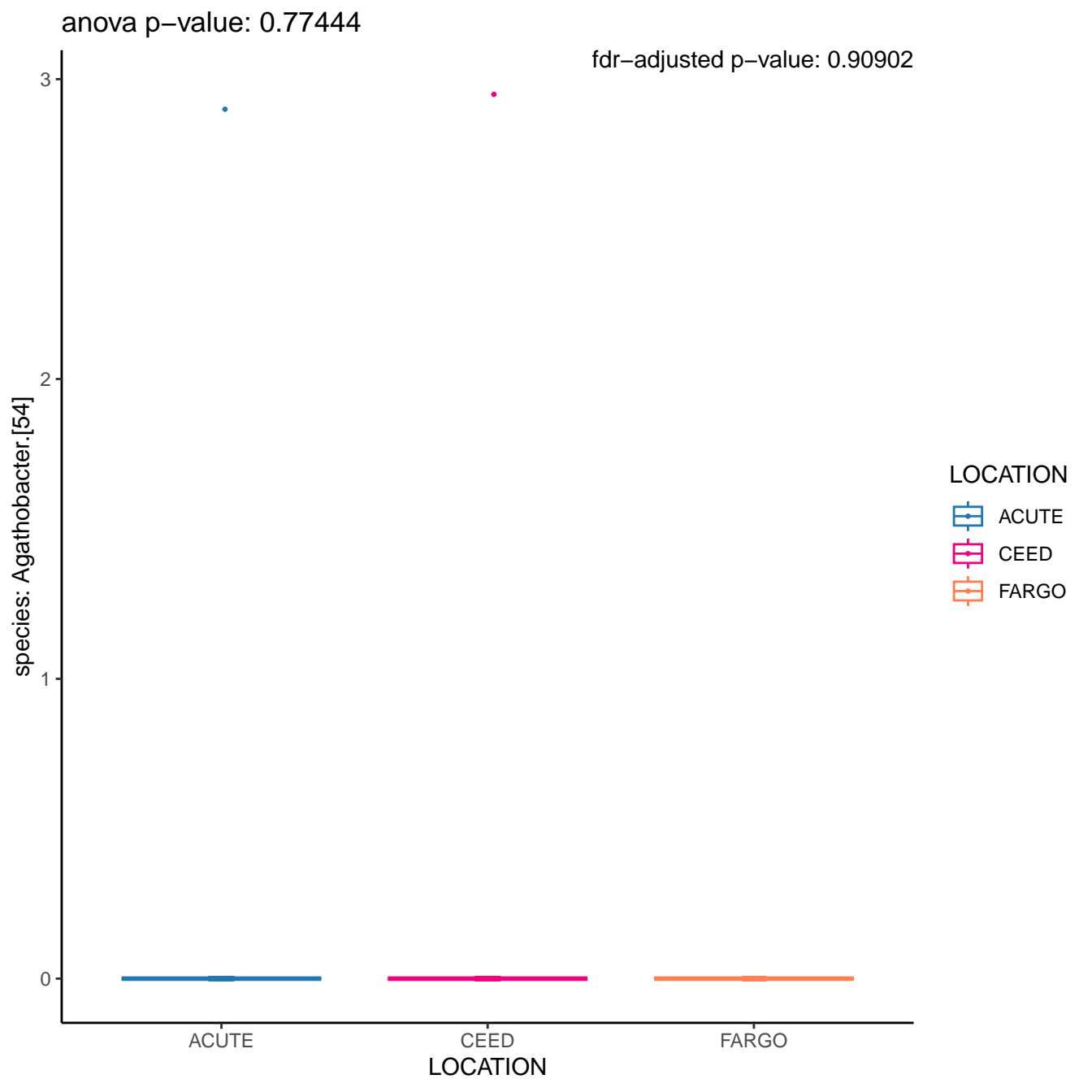
3

ACUTE

CEED

FARGO

LOCATION



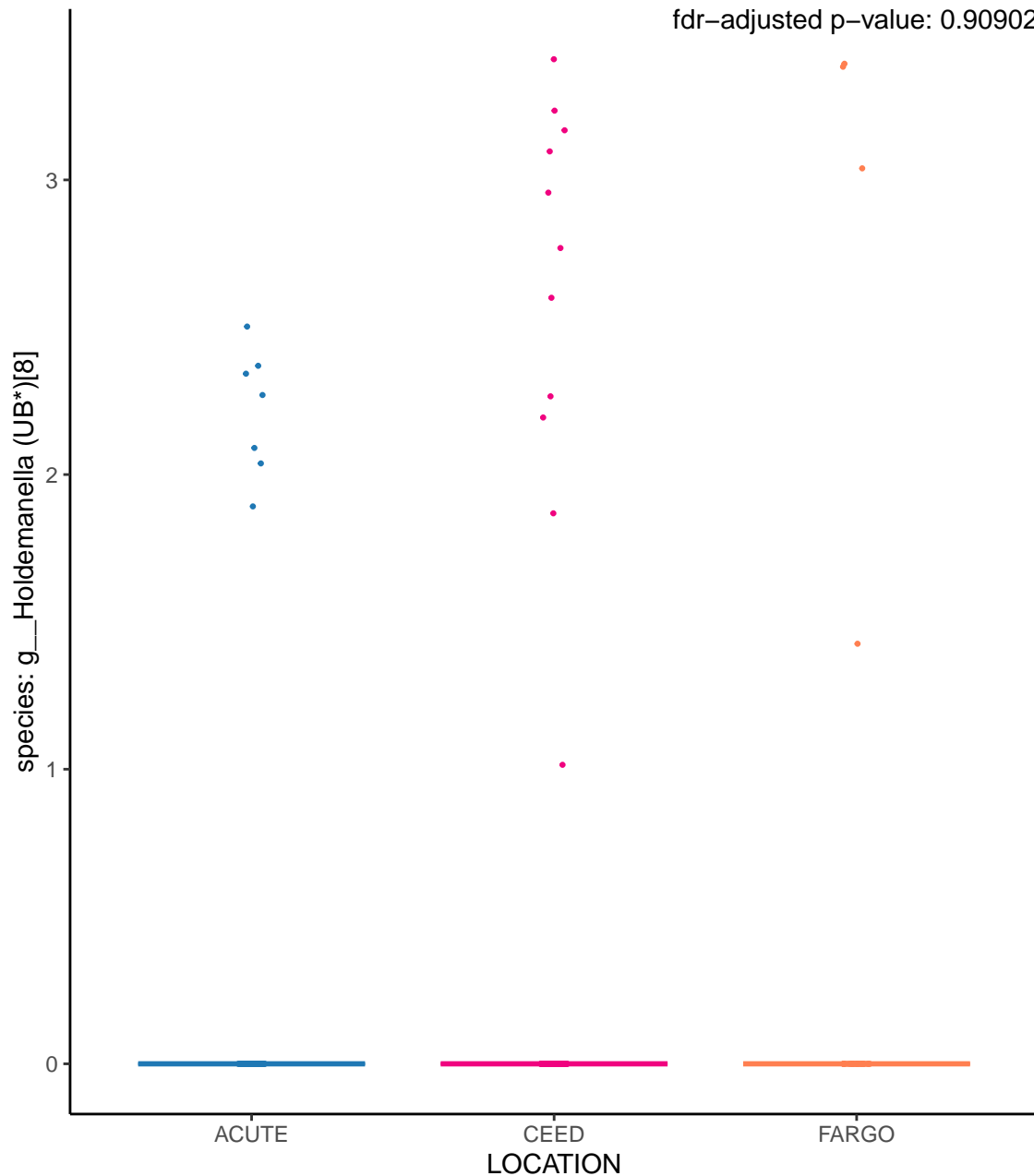
anova p-value: 0.77914

fdr-adjusted p-value: 0.90902

species: g__Holdemanella (UB*)[8]

LOCATION

ACUTE
CEED
FARGO



anova p-value: 0.78218

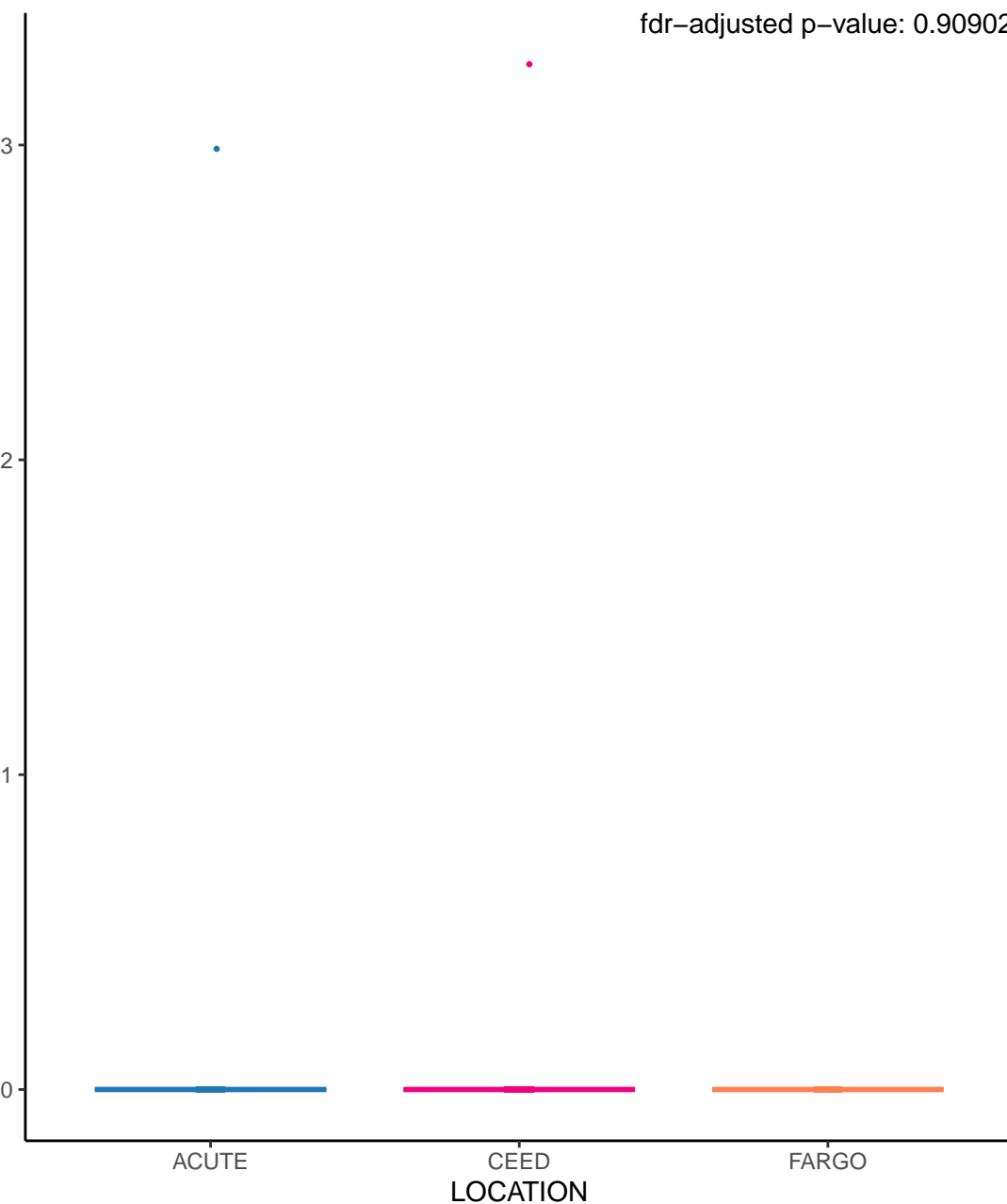
fdr-adjusted p-value: 0.90902

species: UCG.002.[137]

LOCATION

ACUTE
CEED
FARGO

species: UCG.002.[137]

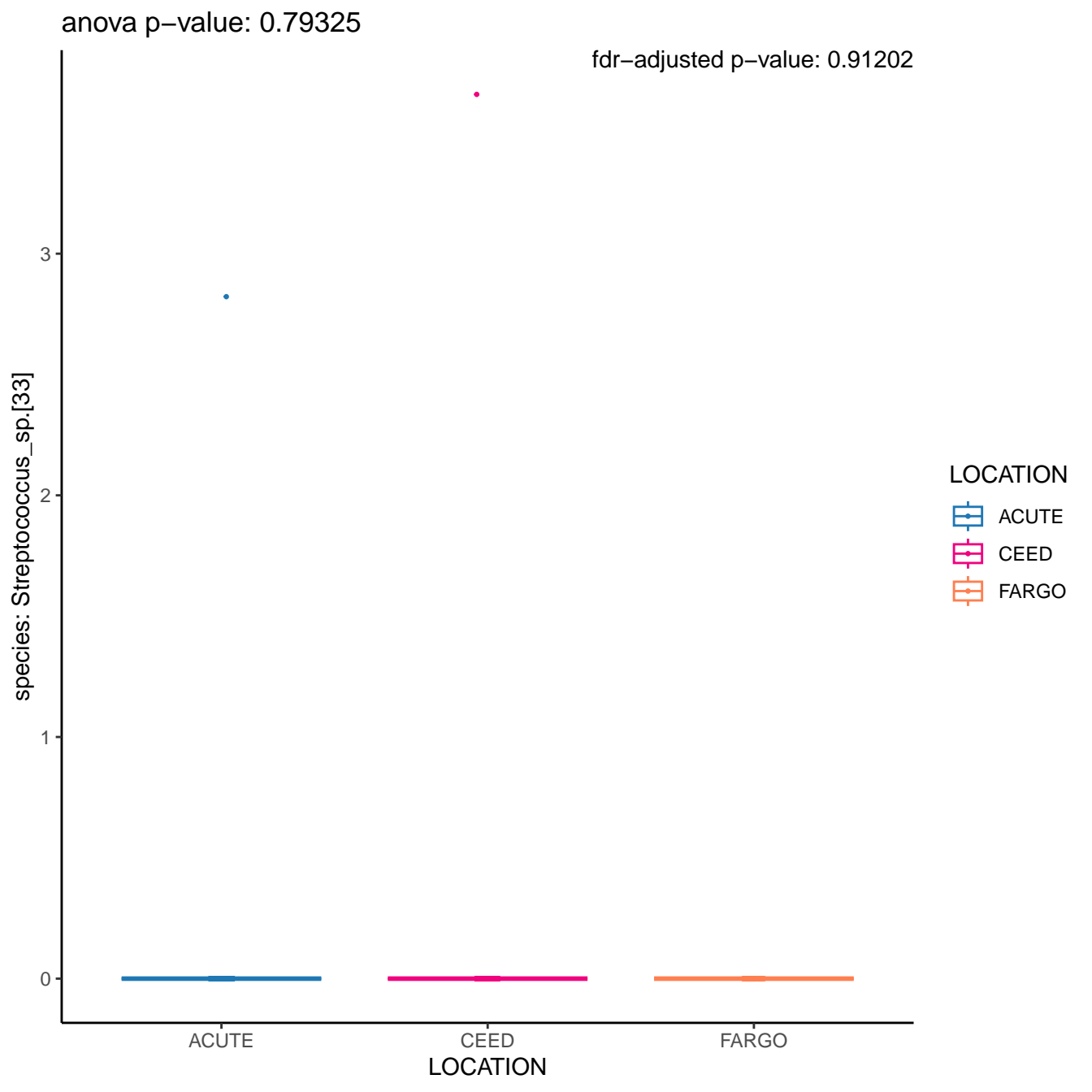
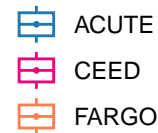


anova p-value: 0.79325

fdr-adjusted p-value: 0.91202

species: Streptococcus_sp.[33]

LOCATION



anova p-value: 0.79536

fdr-adjusted p-value: 0.91202

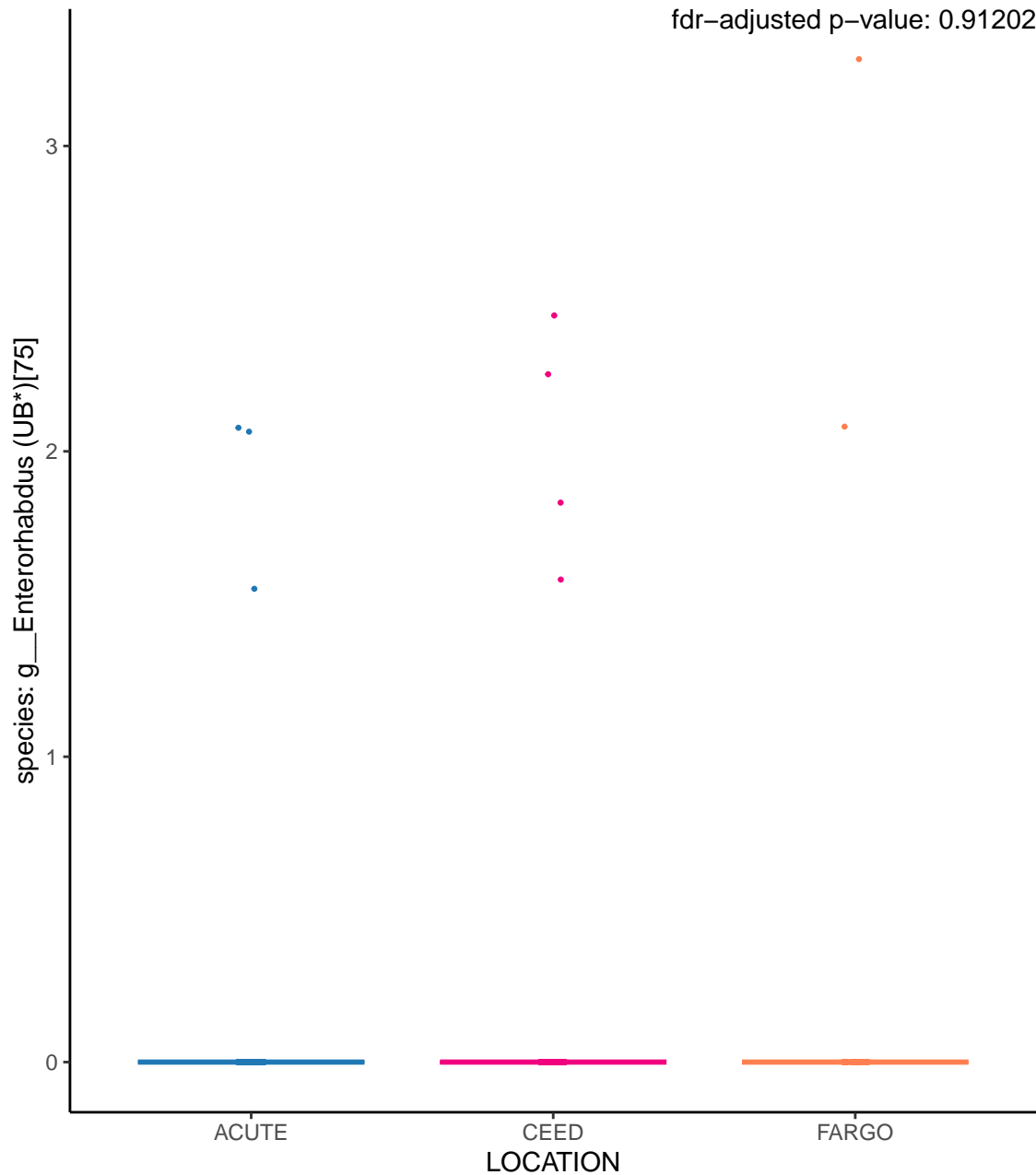
species: g__Enterorhabdus (UB*)[75]

LOCATION

ACUTE

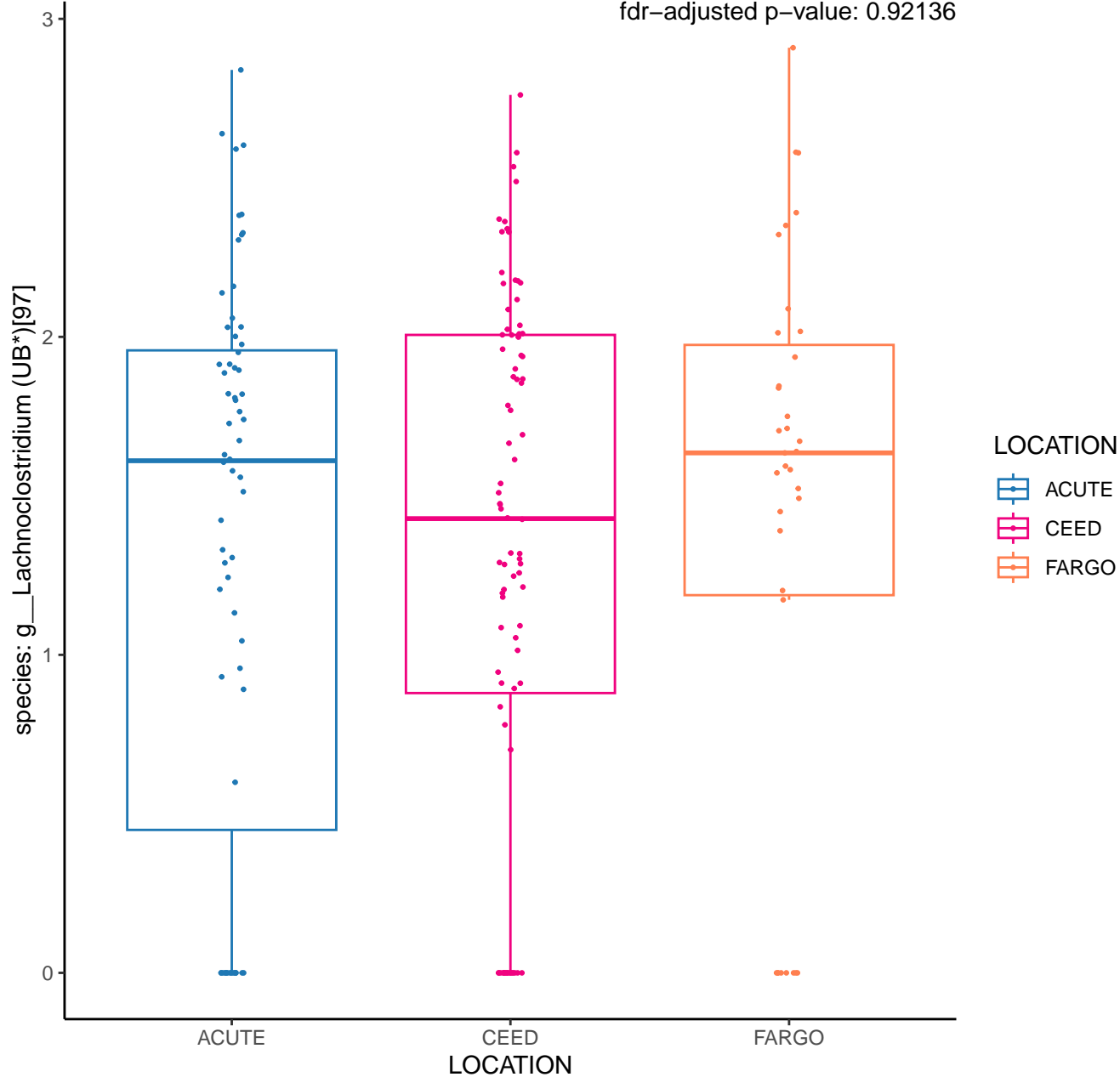
CEED

FARGO



anova p-value: 0.80887

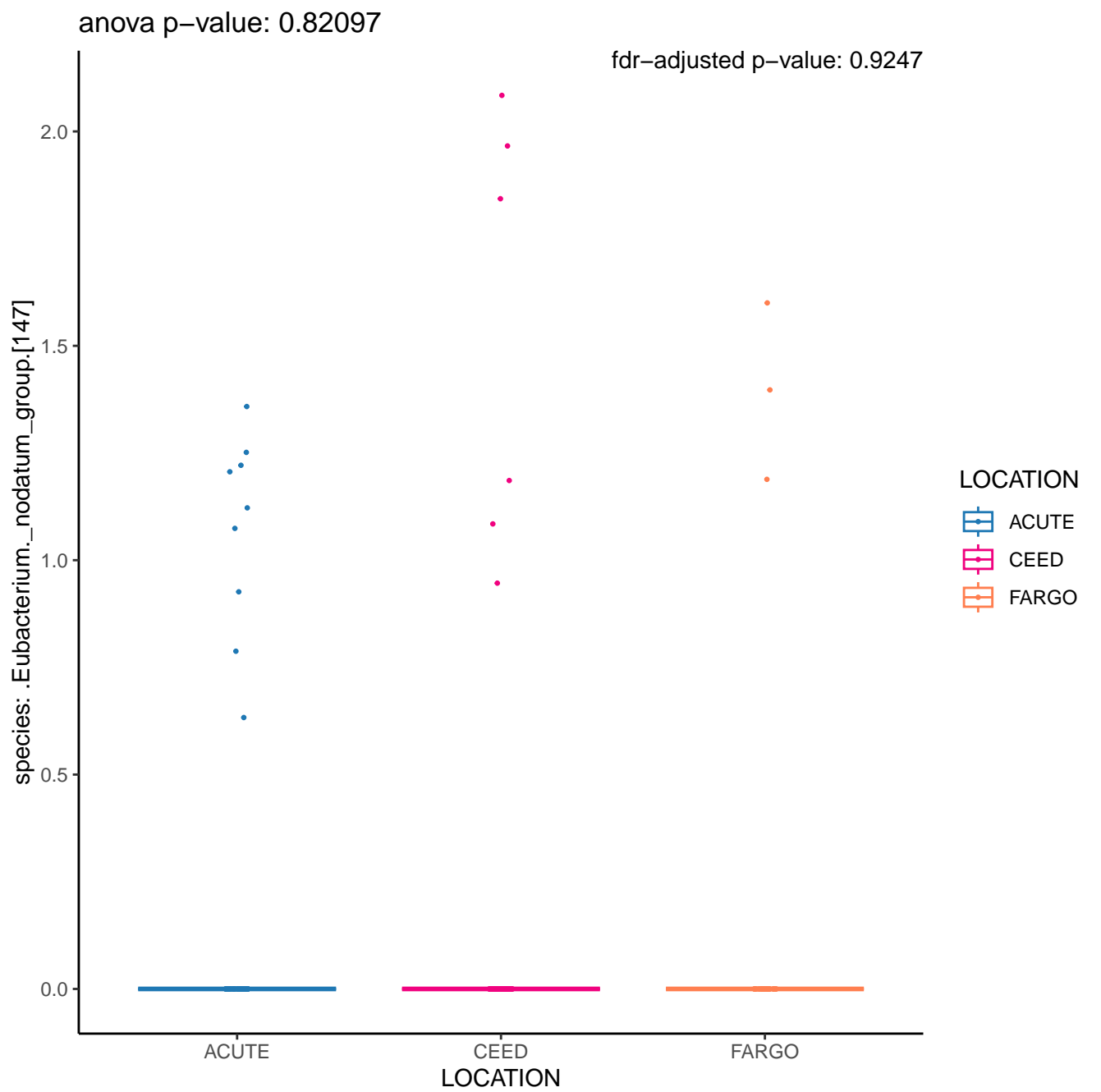
fdr-adjusted p-value: 0.92136



anova p-value: 0.82097

fdr-adjusted p-value: 0.9247

species: .Eubacterium._nodatum_group.[147]

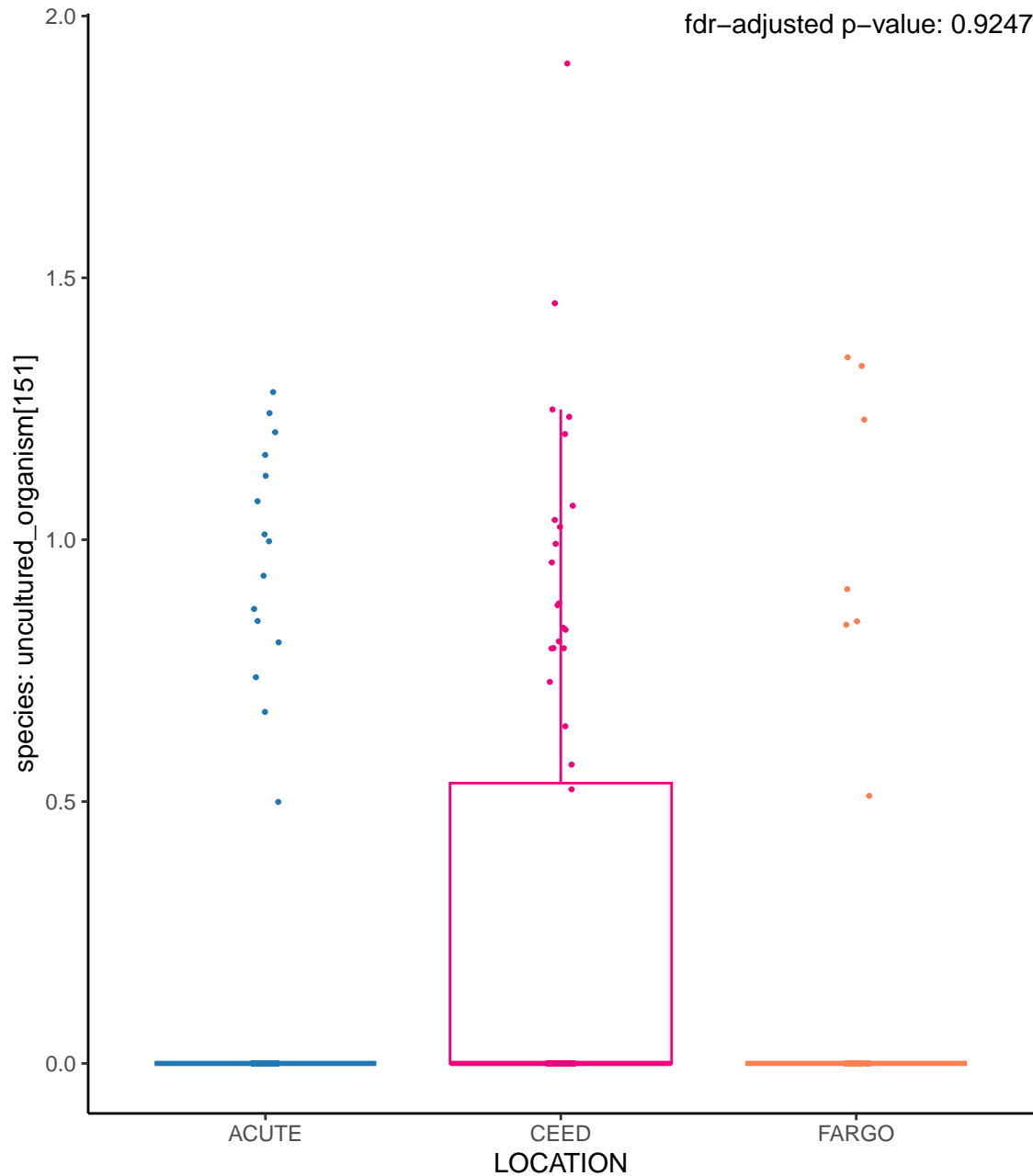


anova p-value: 0.82977

fdr-adjusted p-value: 0.9247

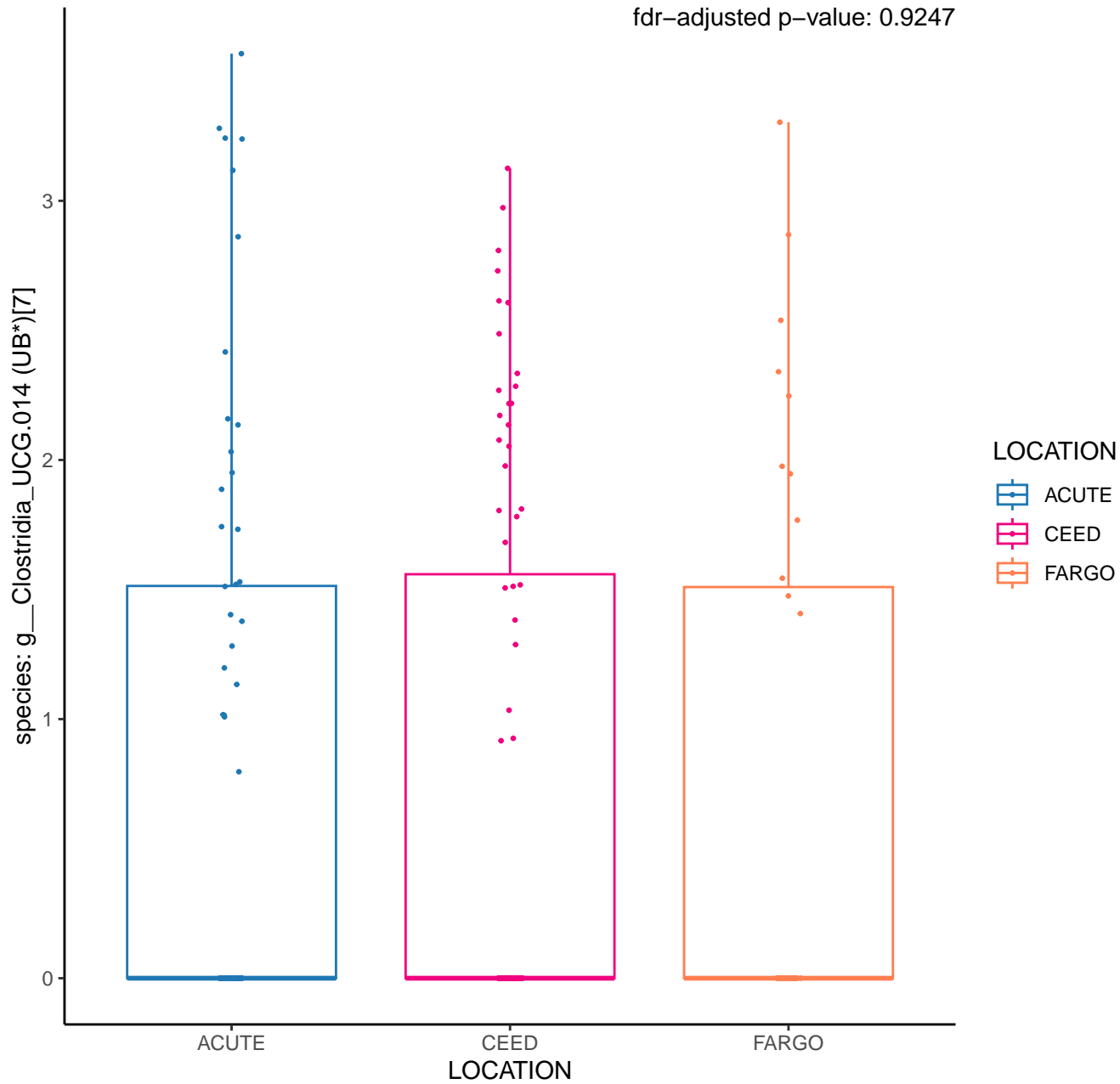
species: uncultured_organism[151]

LOCATION



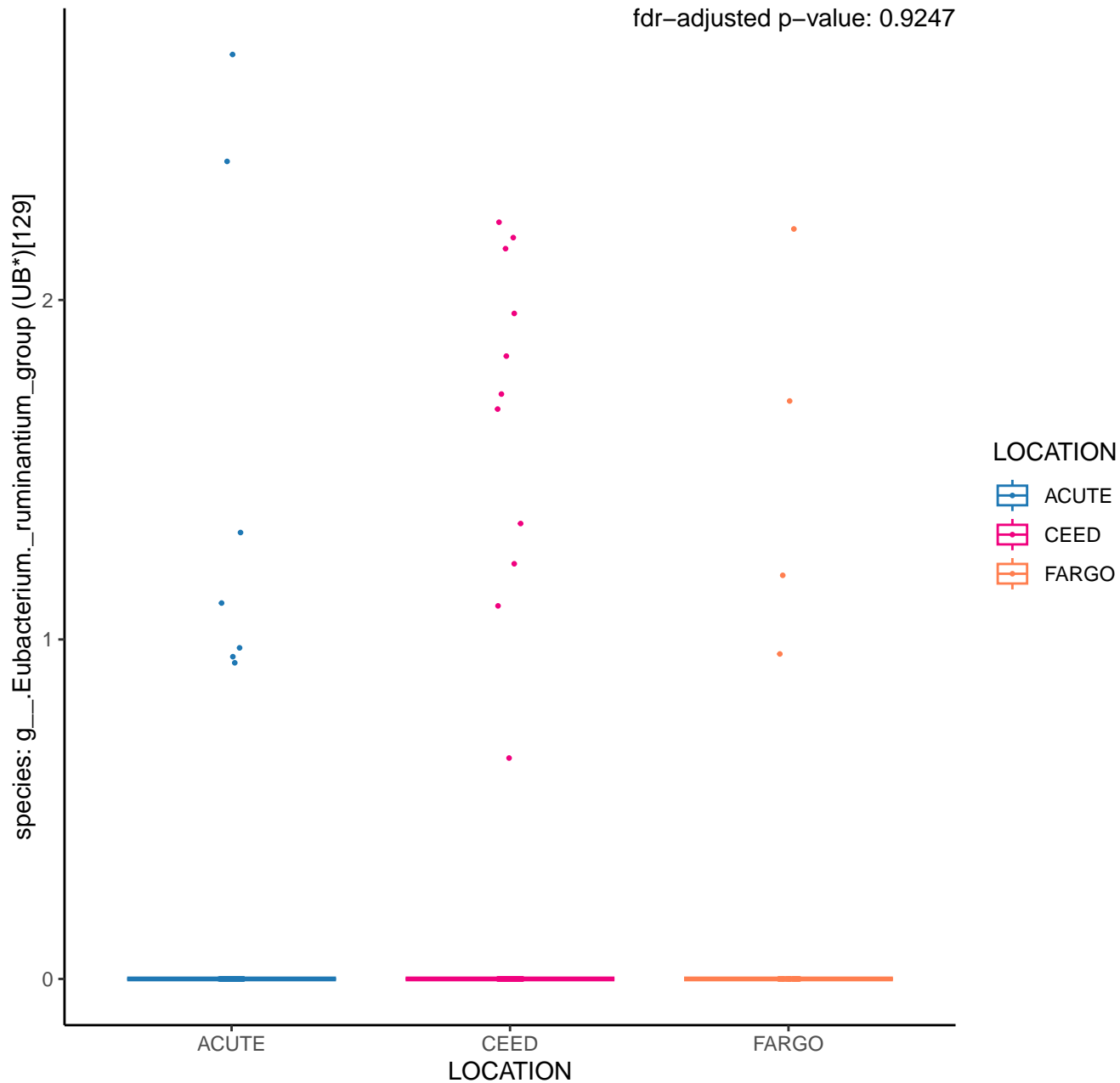
anova p-value: 0.83073

fdr-adjusted p-value: 0.9247



anova p-value: 0.8333

fdr-adjusted p-value: 0.9247



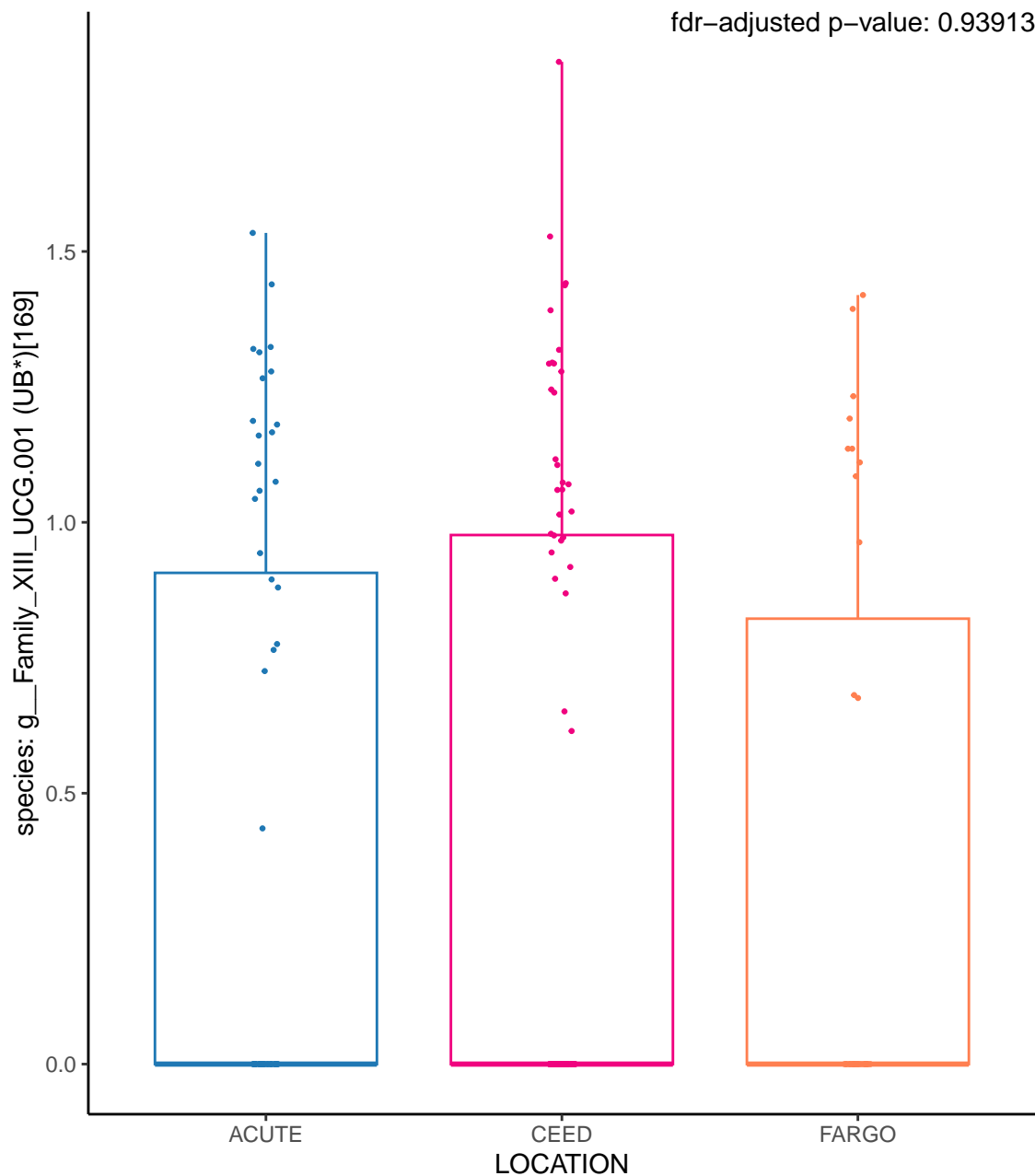
anova p-value: 0.85304

fdr-adjusted p-value: 0.93913

species: g__Family_XIII_UCG.001 (UB*)[169]

LOCATION

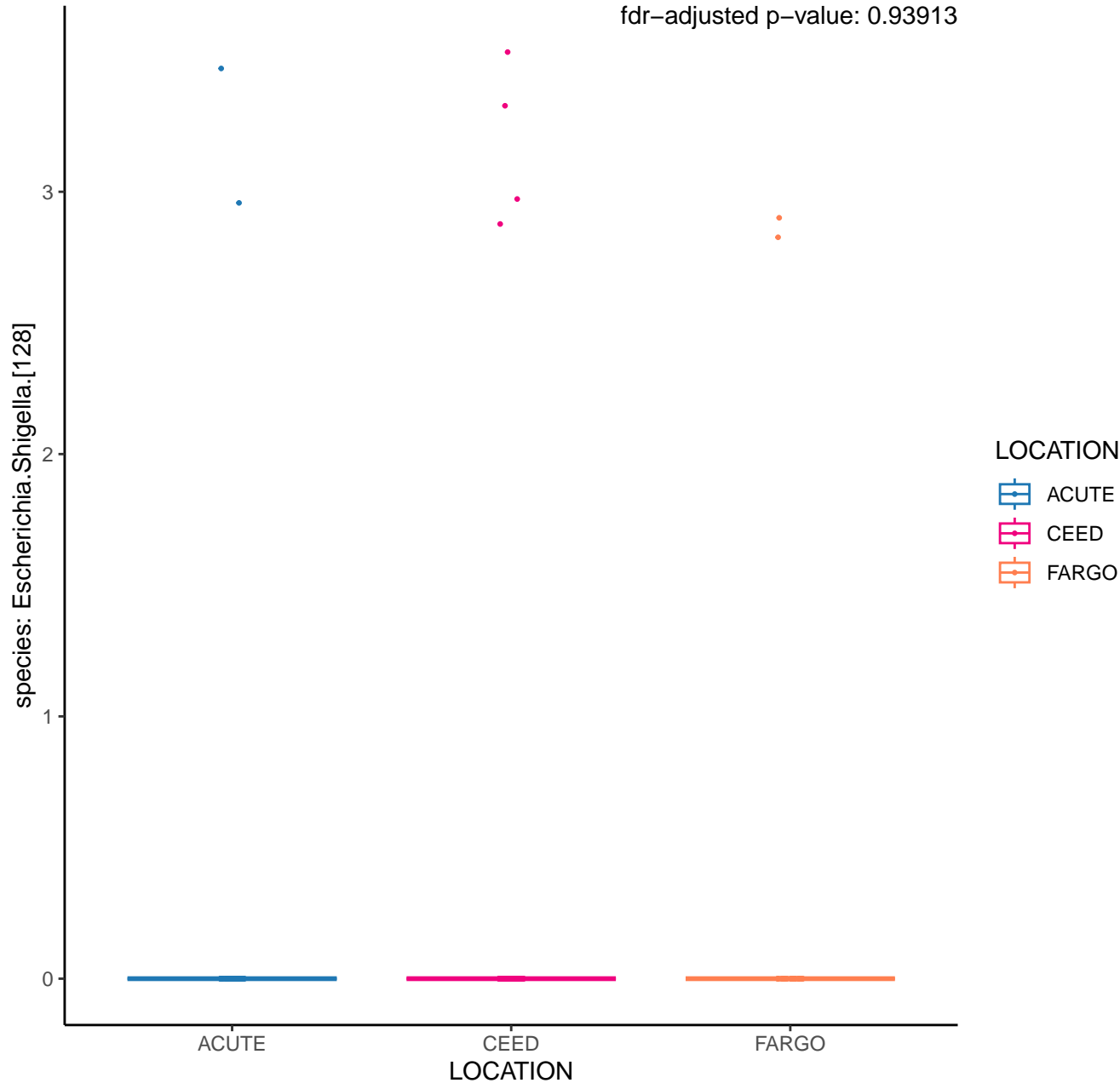
ACUTE
CEED
FARGO



anova p-value: 0.85723

fdr-adjusted p-value: 0.93913

species: *Escherichia.Shigella*. [128]

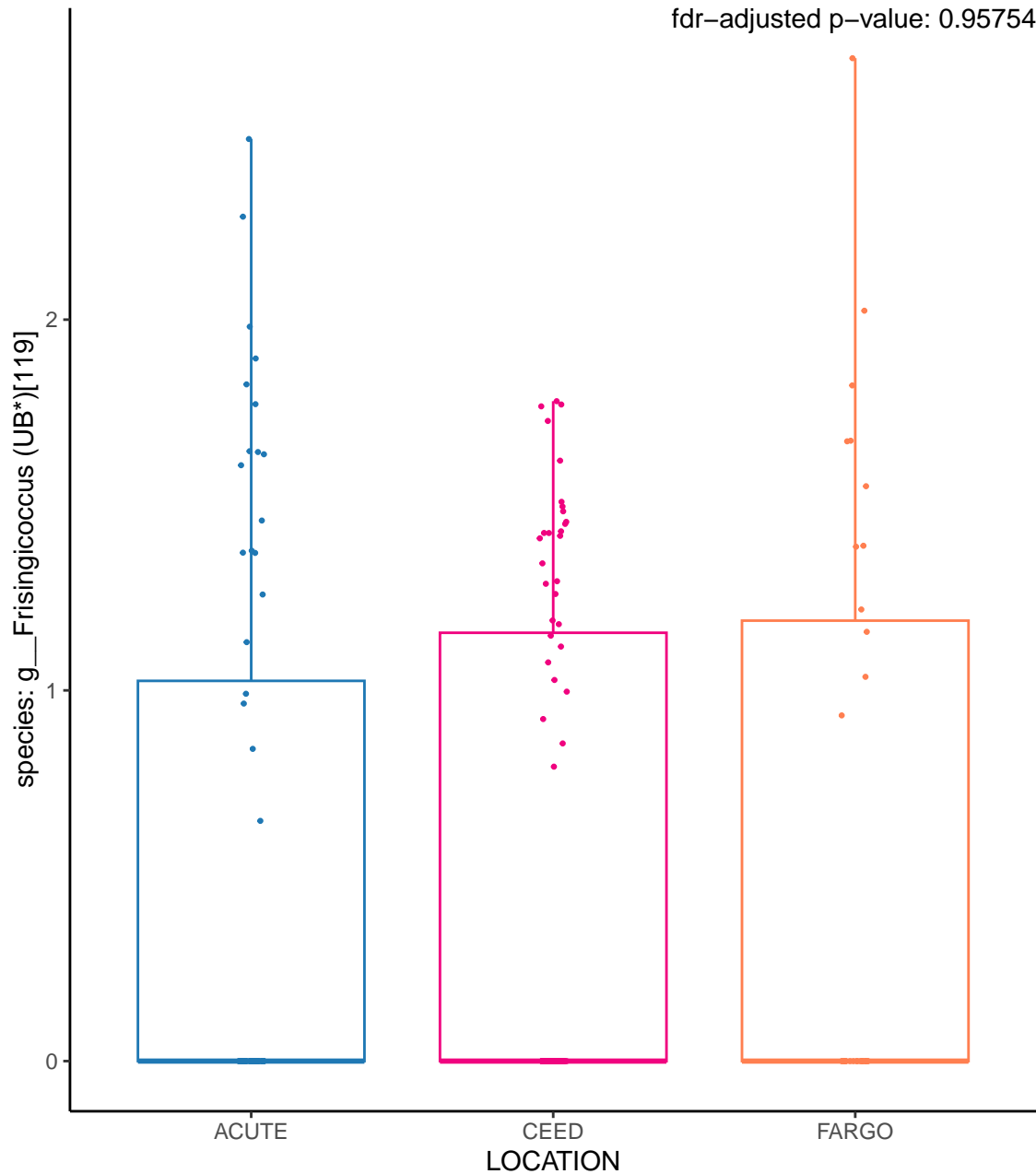


anova p-value: 0.88483

fdr-adjusted p-value: 0.95754

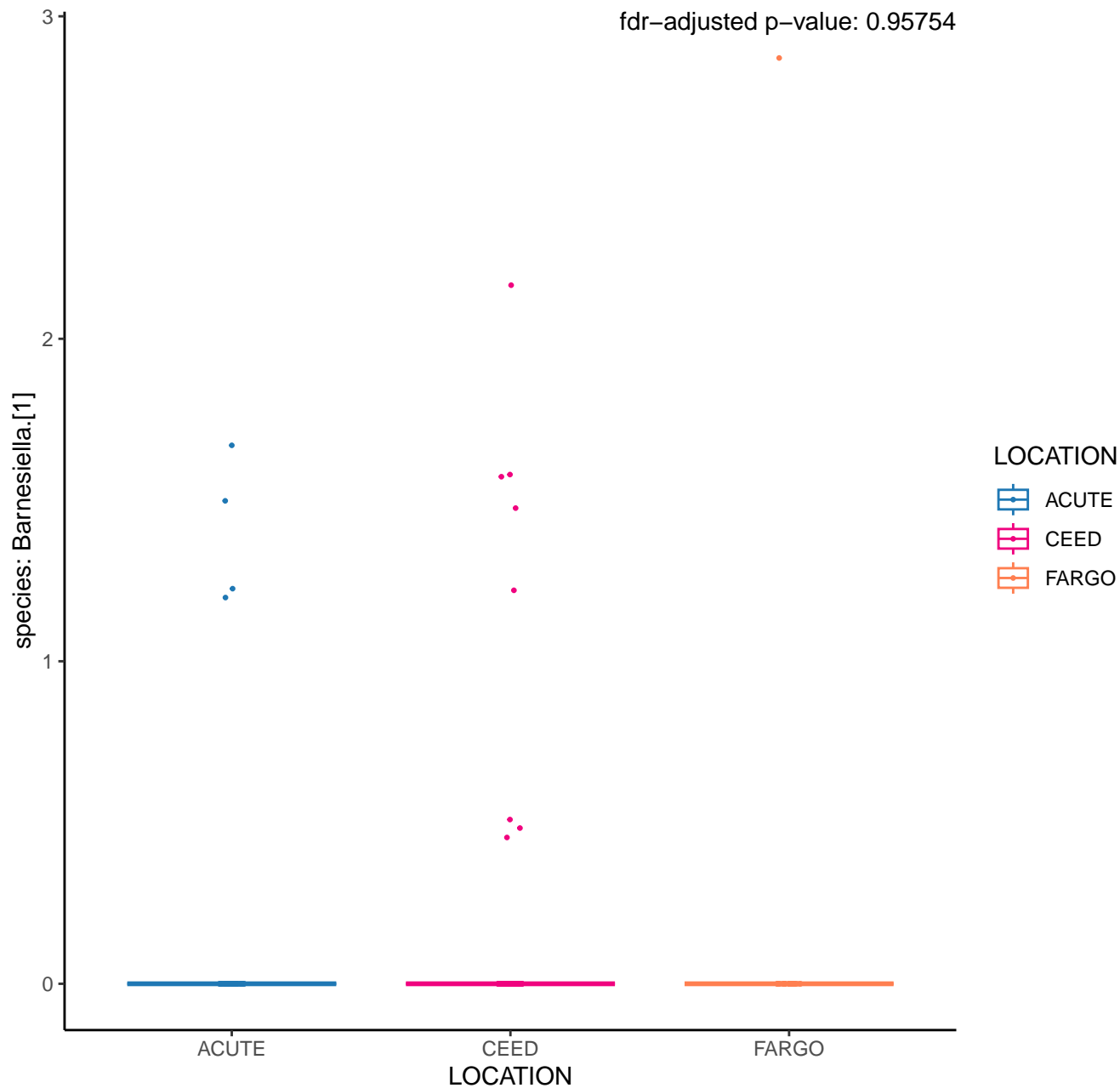
species: g__Frisingicoccus (UB*)[119]

LOCATION

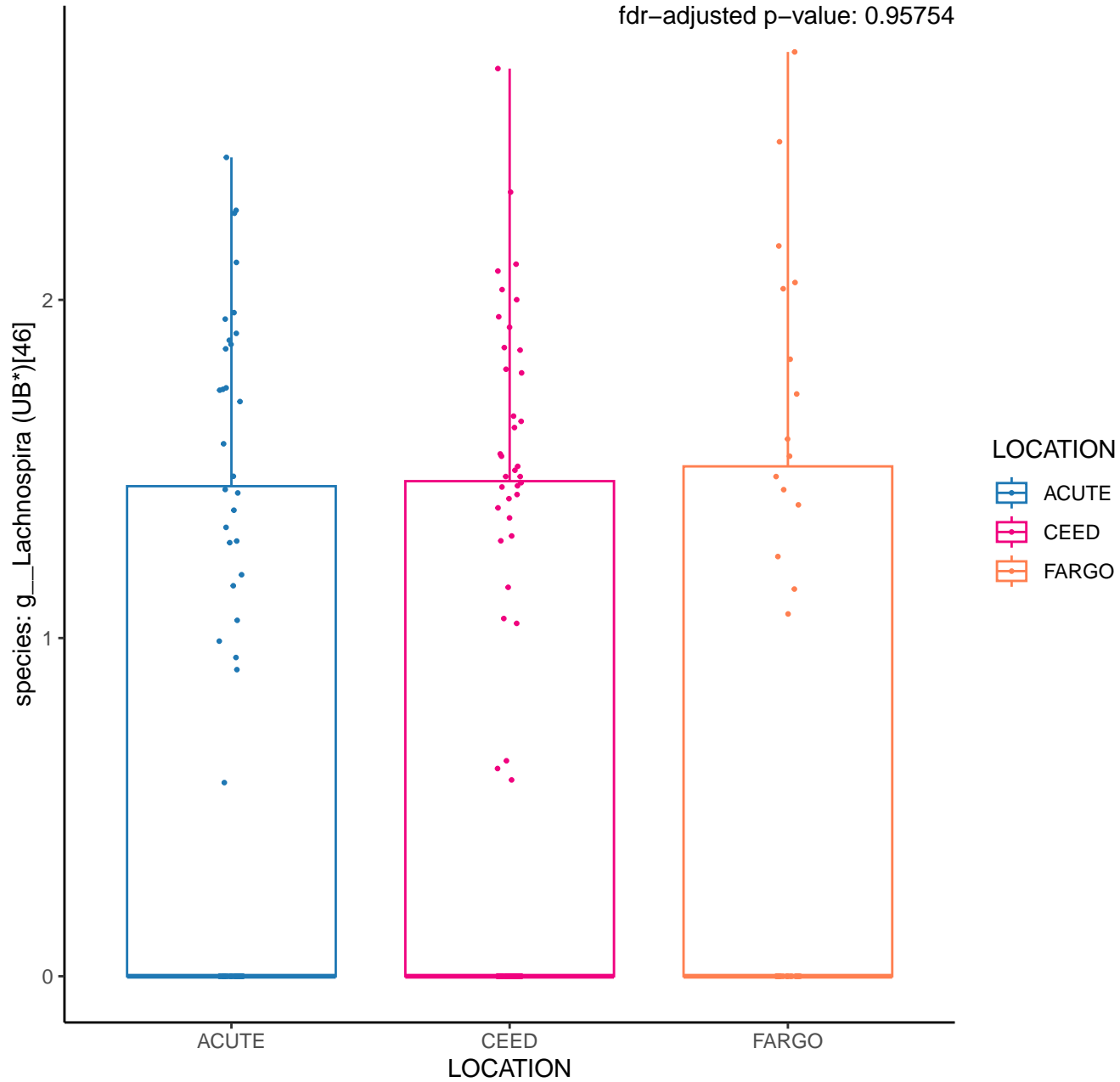


anova p-value: 0.89774

fdr-adjusted p-value: 0.95754

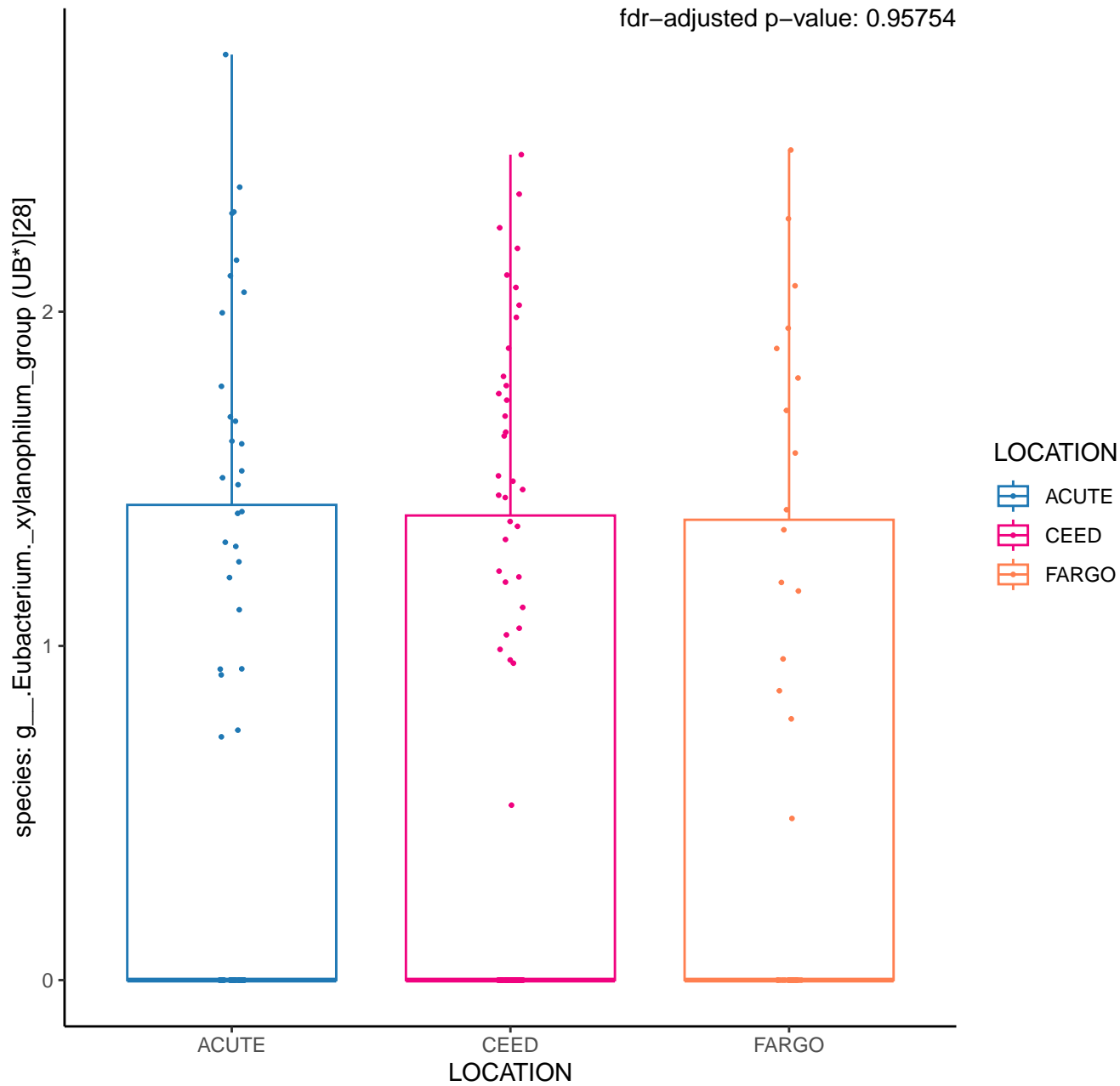


fdr-adjusted p-value: 0.95754



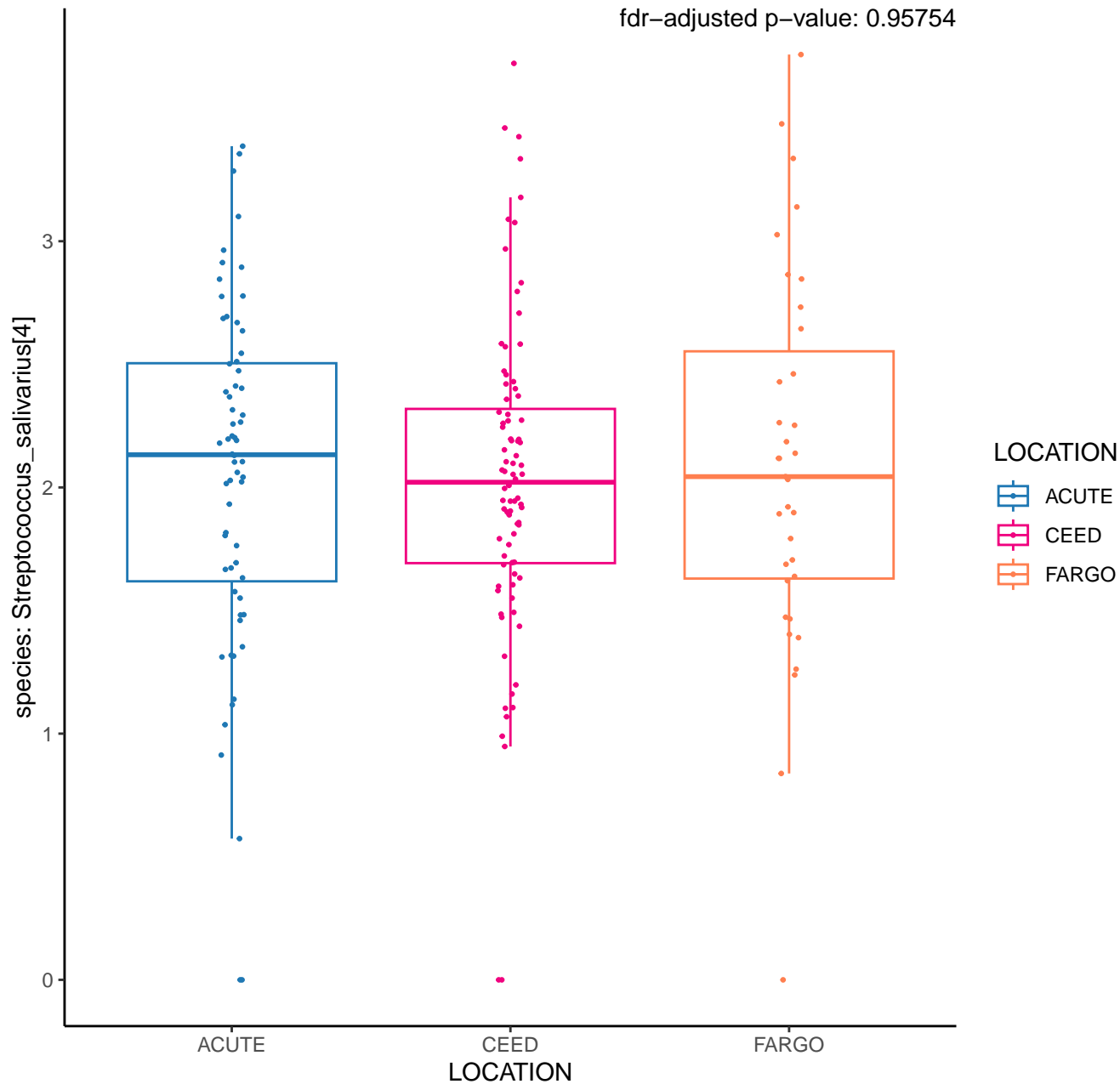
anova p-value: 0.90177

fdr-adjusted p-value: 0.95754



anova p-value: 0.90187

fdr-adjusted p-value: 0.95754



anova p-value: 0.91838

fdr-adjusted p-value: 0.96909

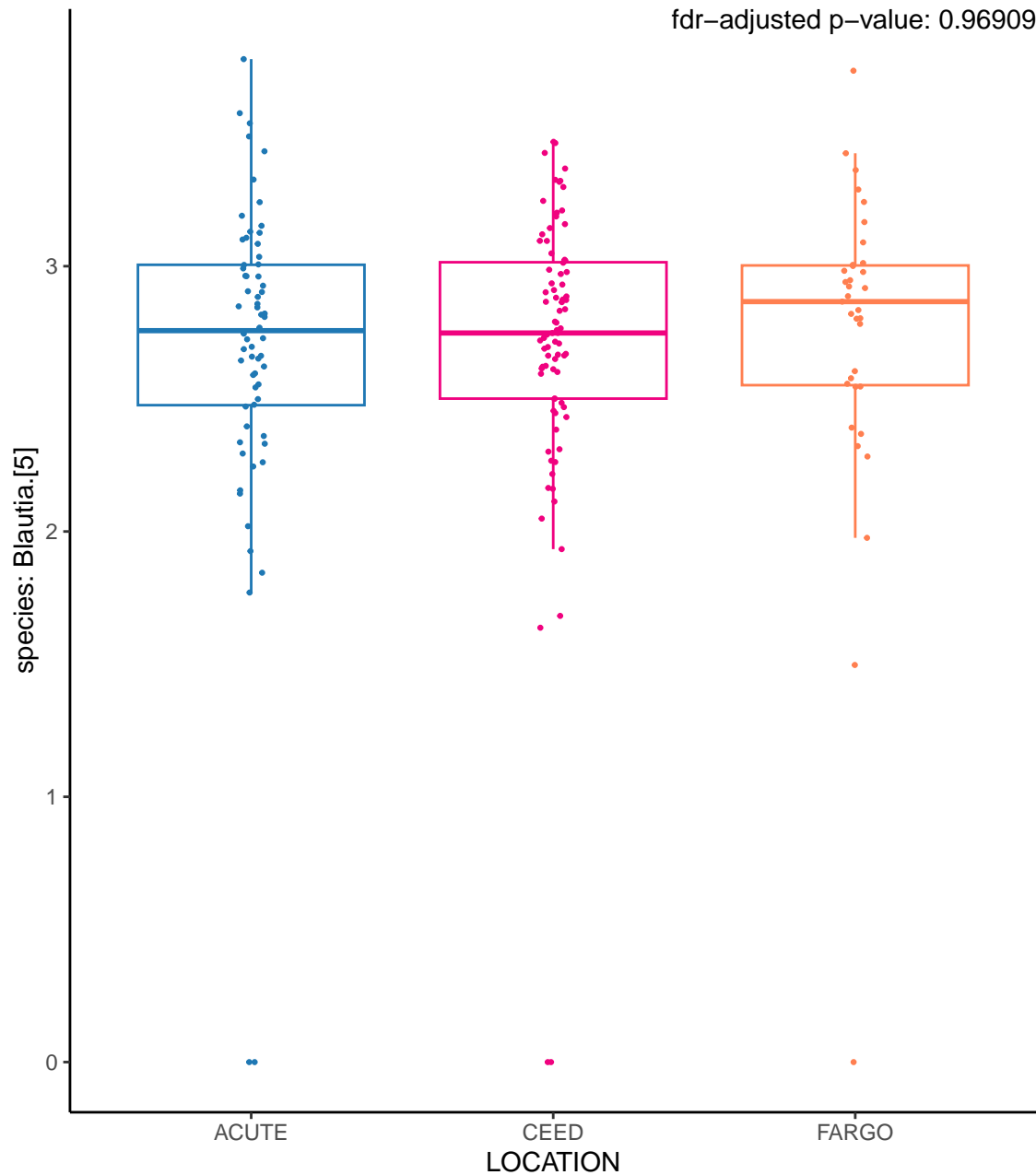
species: *Blautia*.[5]

LOCATION

ACUTE

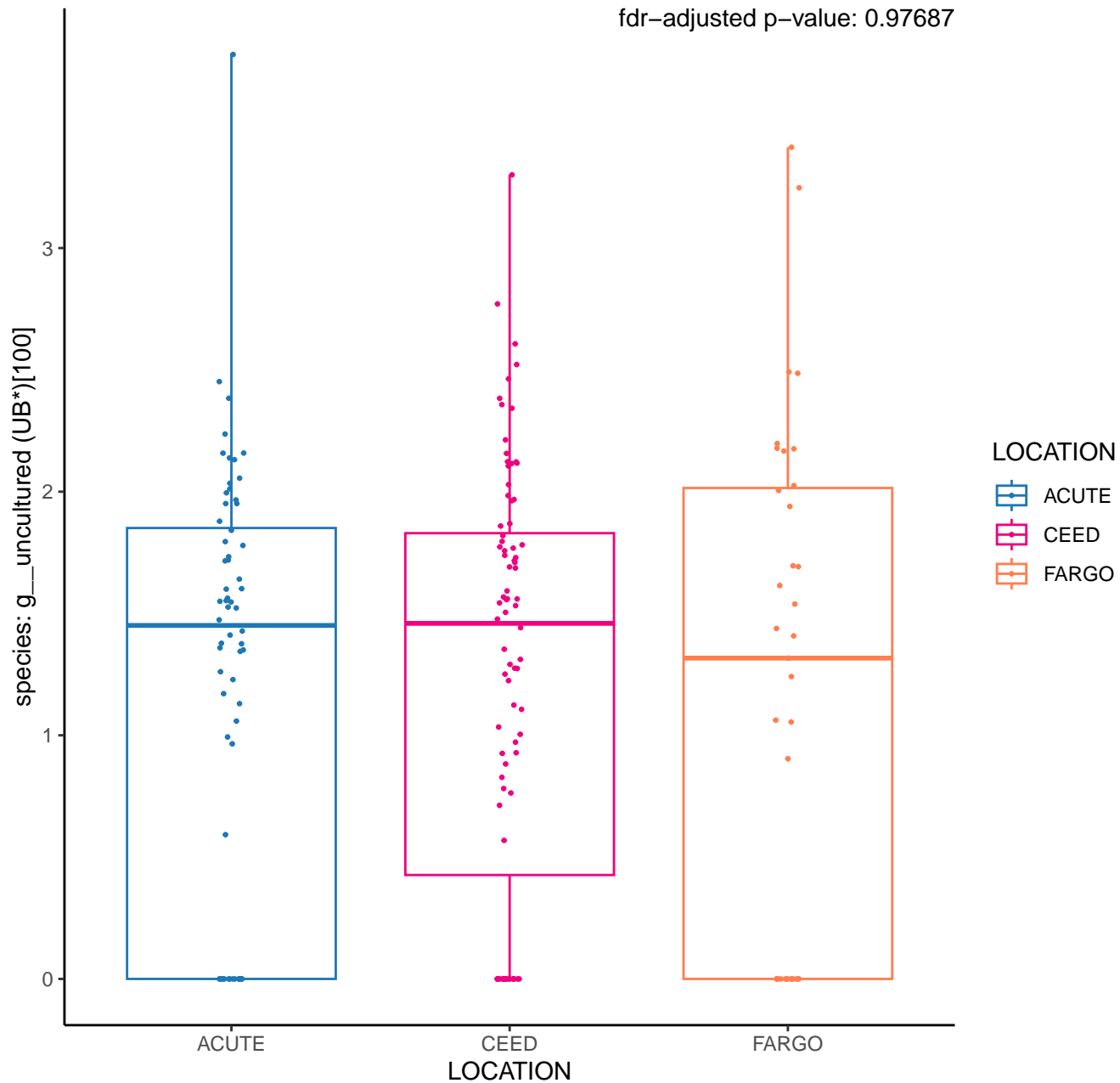
CEED

FARGO



anova p-value: 0.93566

fdr-adjusted p-value: 0.97687

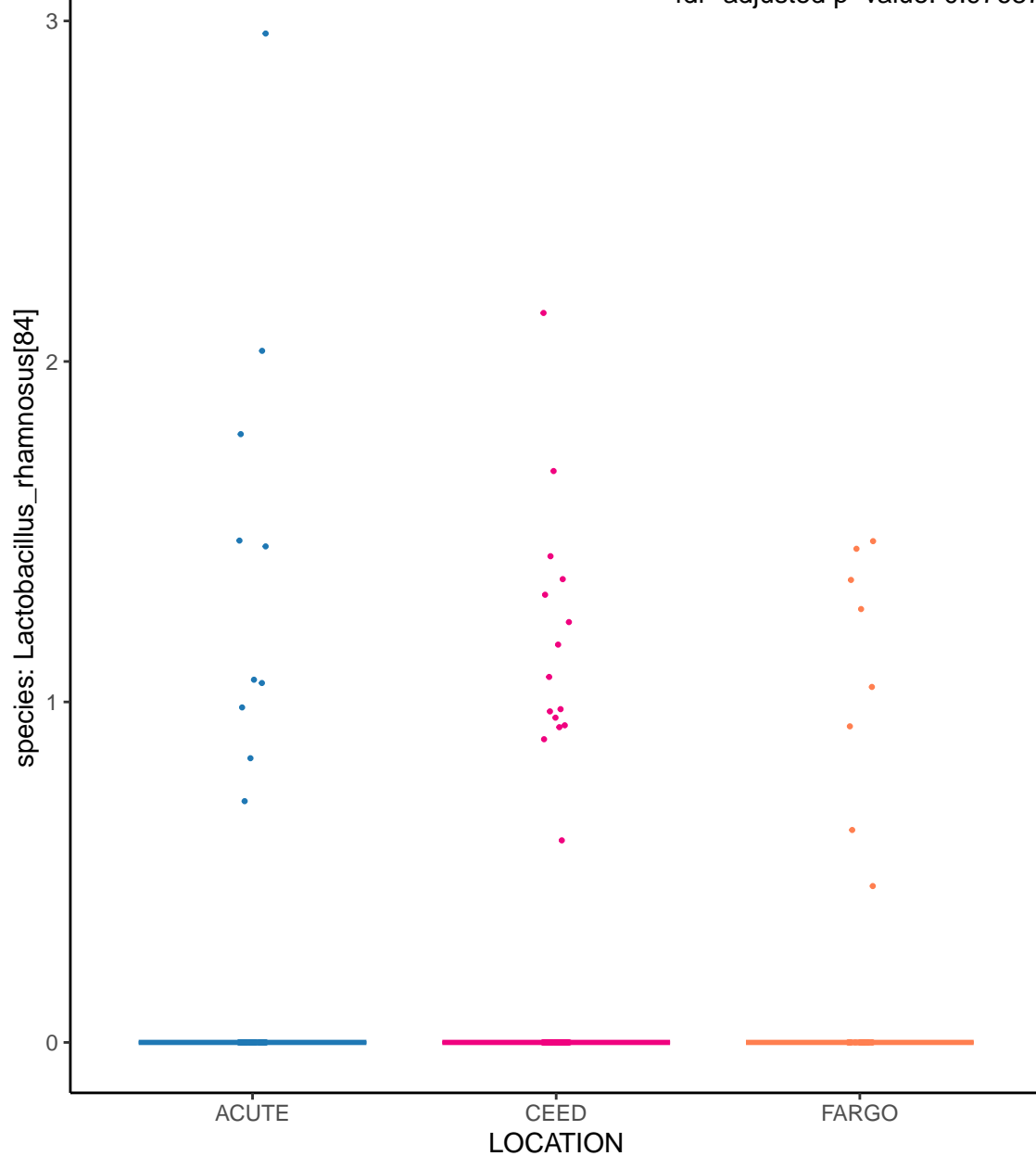
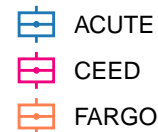


anova p-value: 0.942

fdr-adjusted p-value: 0.97687

species: *Lactobacillus_rhamnosus*[84]

LOCATION



anova p-value: 0.94279

fdr-adjusted p-value: 0.97687

species: Enterobacter.[126]

LOCATION

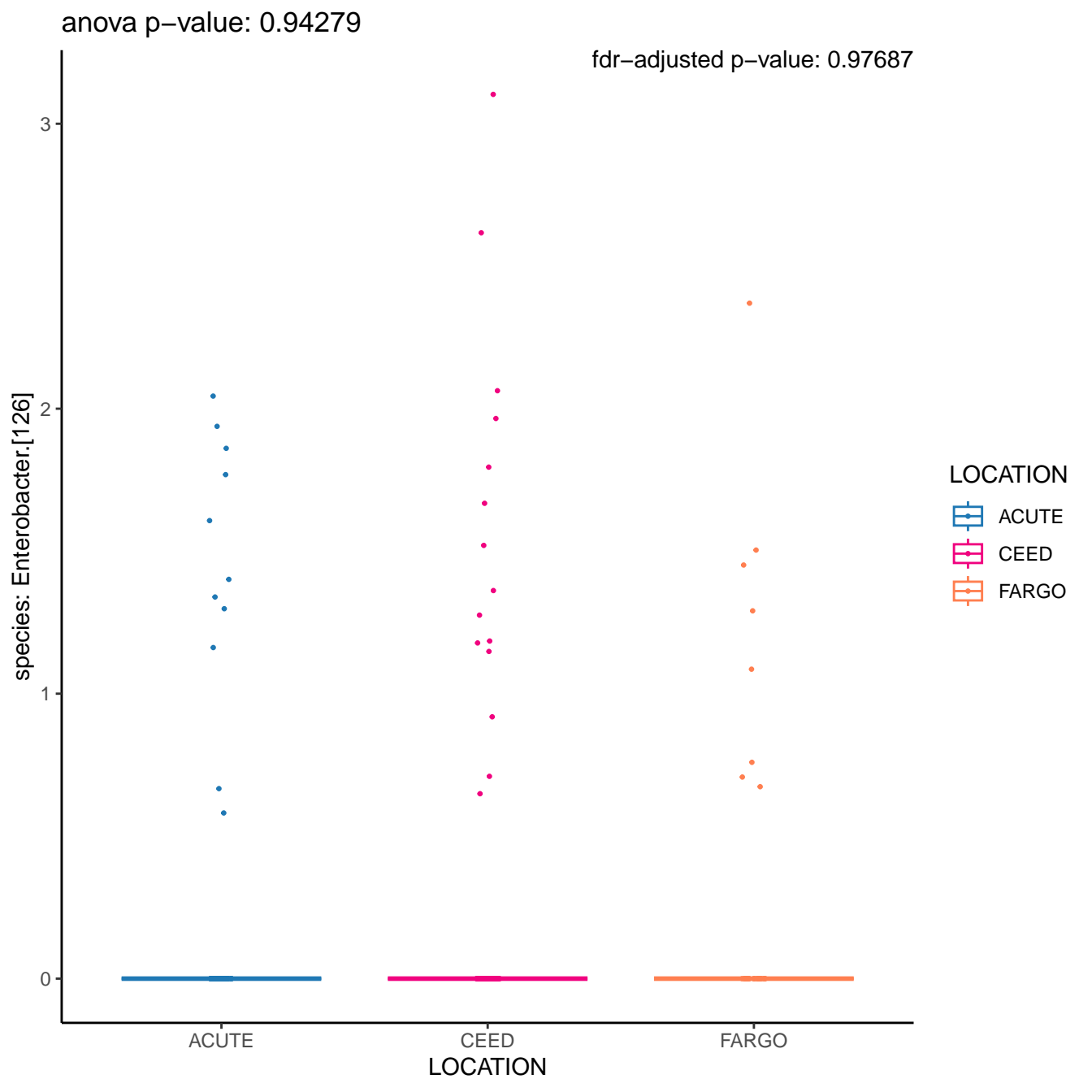
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CEED

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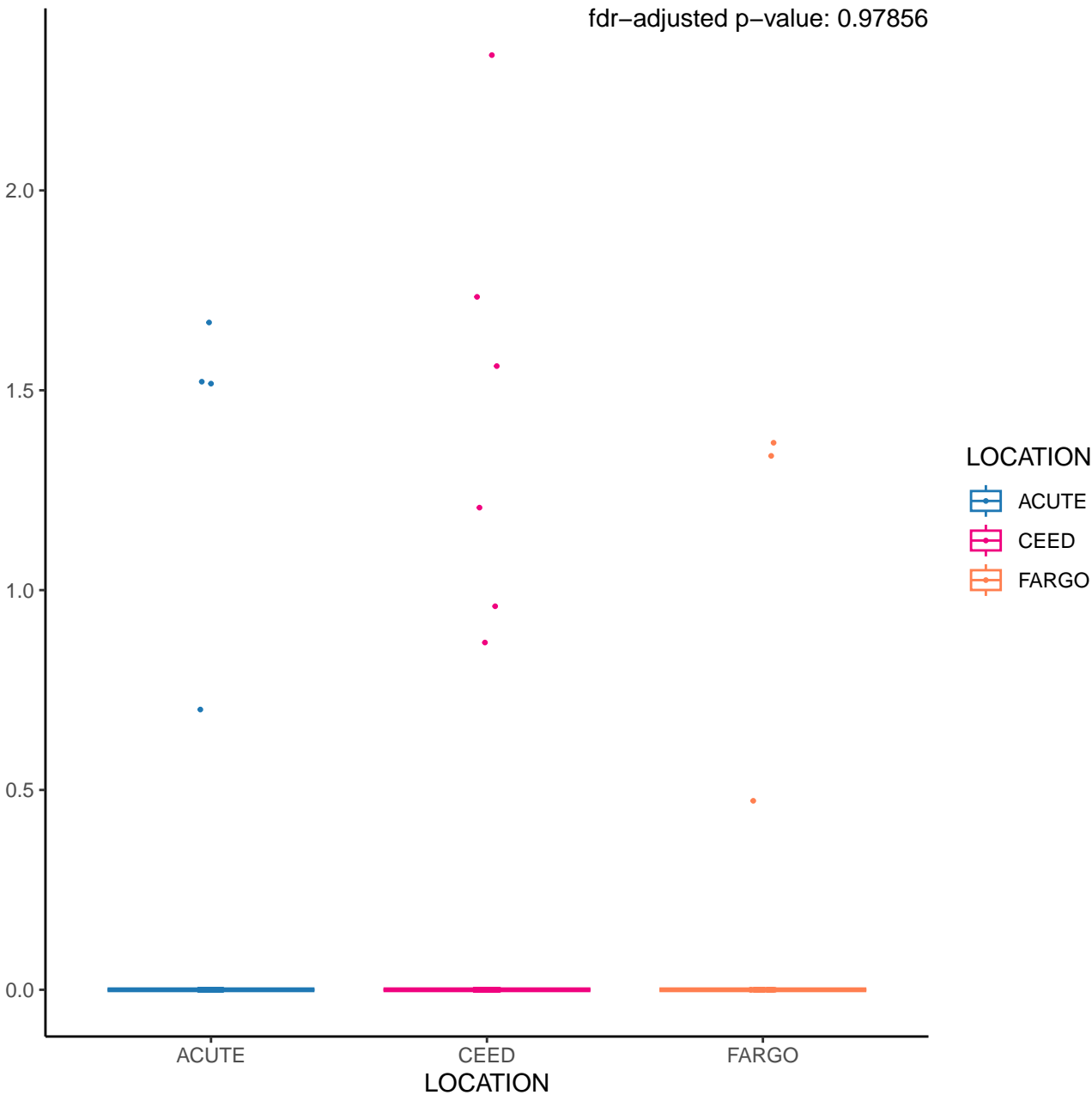
LOCATION



anova p-value: 0.95208

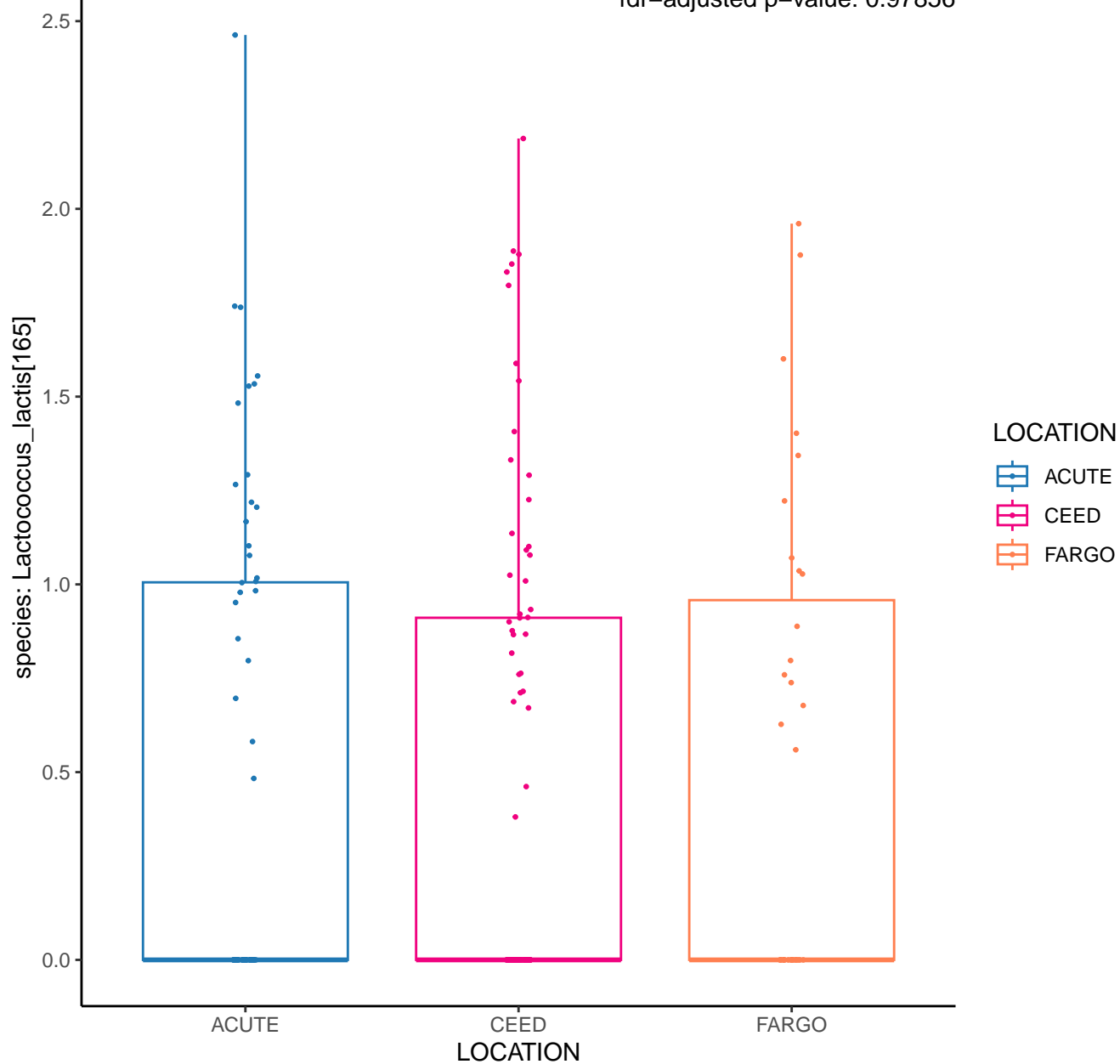
fdr-adjusted p-value: 0.97856

species: g__Sutterella (UB*)[160]



anova p-value: 0.9558

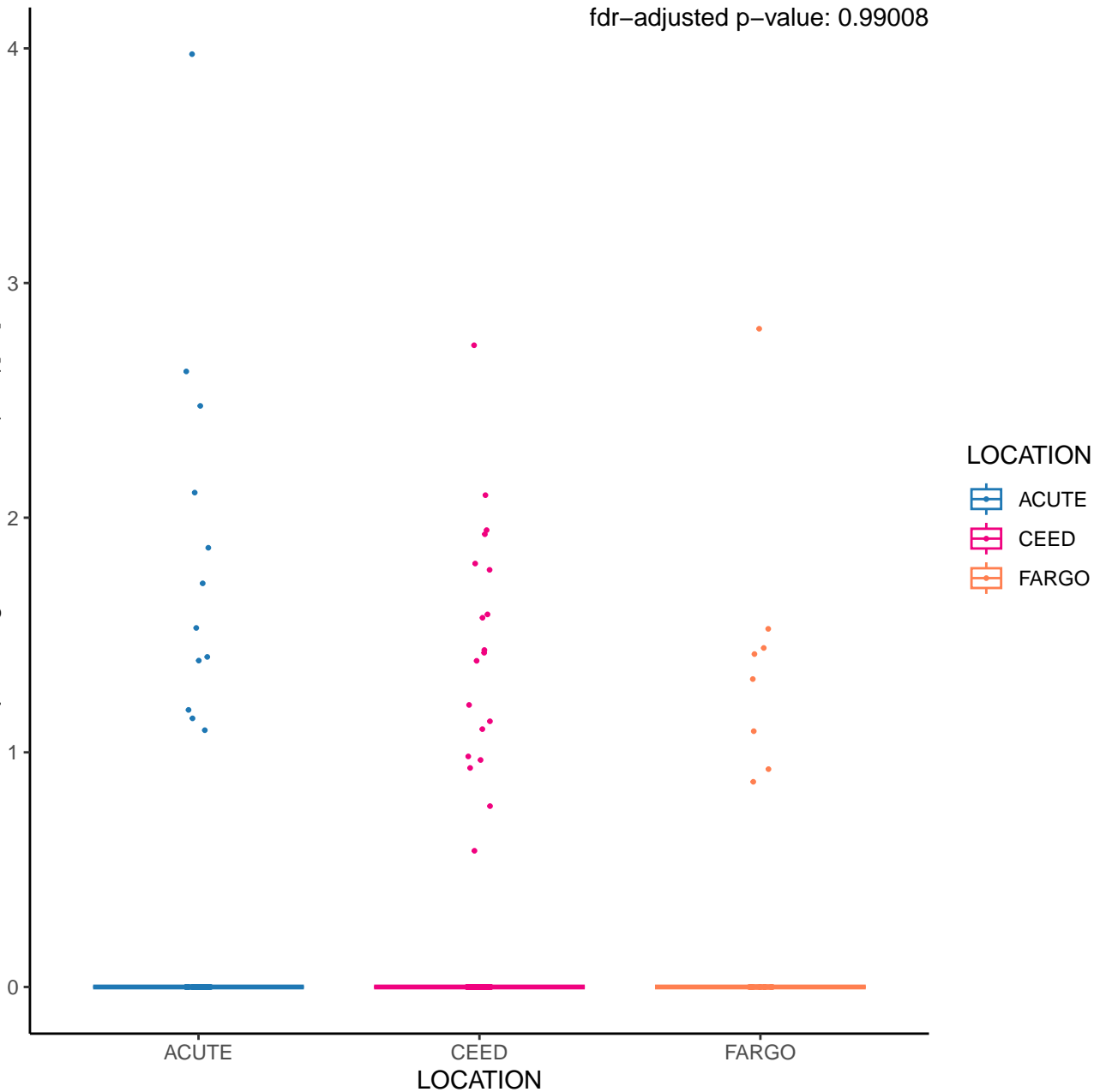
fdr-adjusted p-value: 0.97856



anova p-value: 0.97281

fdr-adjusted p-value: 0.99008

species: g__Lactobacillus (UB*)[91]



anova p-value: 0.98955

fdr-adjusted p-value: 0.99536

species: Family_XIII_AD3011_group.[64]

LOCATION

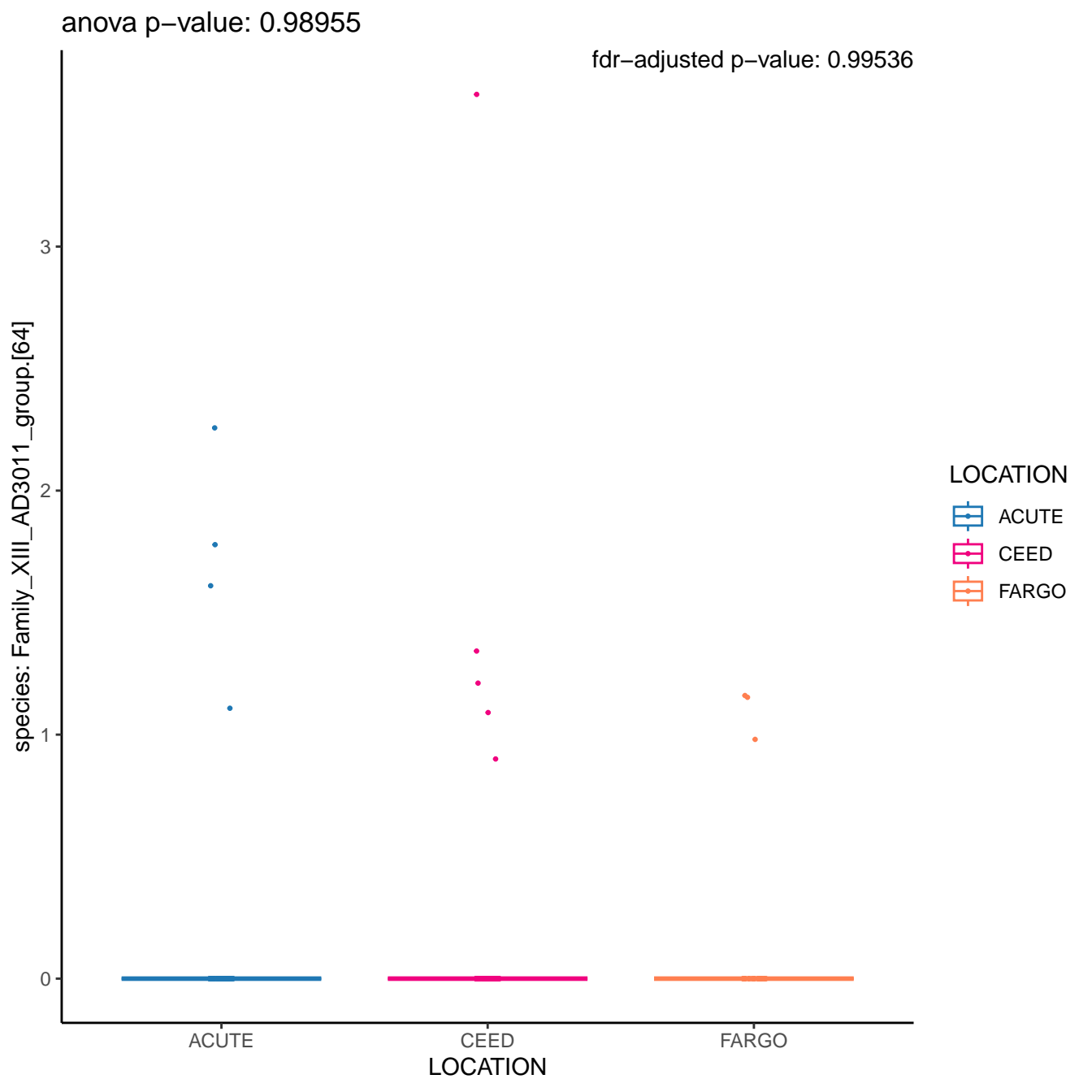
ACUTE
CEED
FARGO

ACUTE

CEED

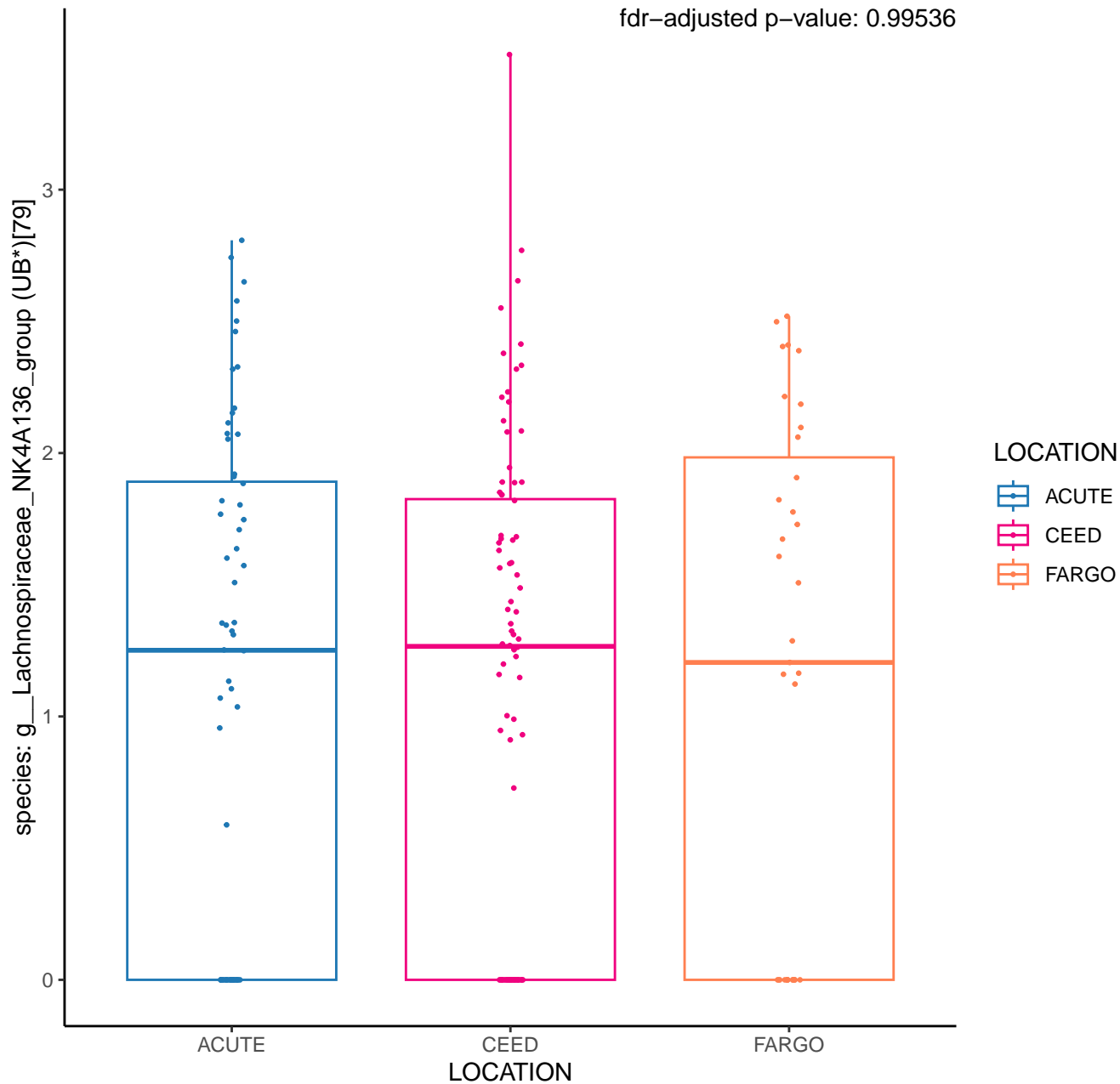
FARGO

LOCATION



anova p-value: 0.98958

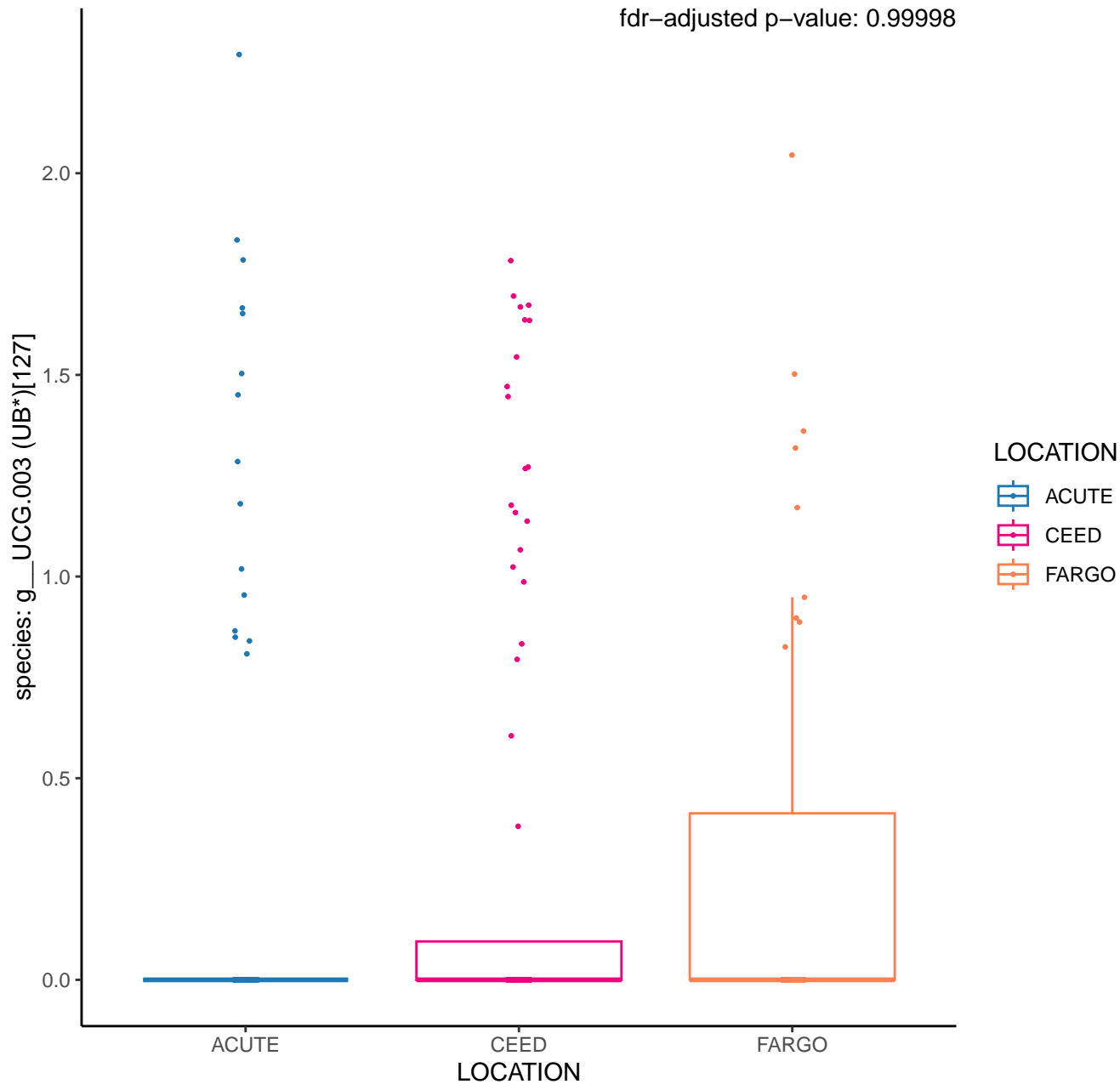
fdr-adjusted p-value: 0.99536



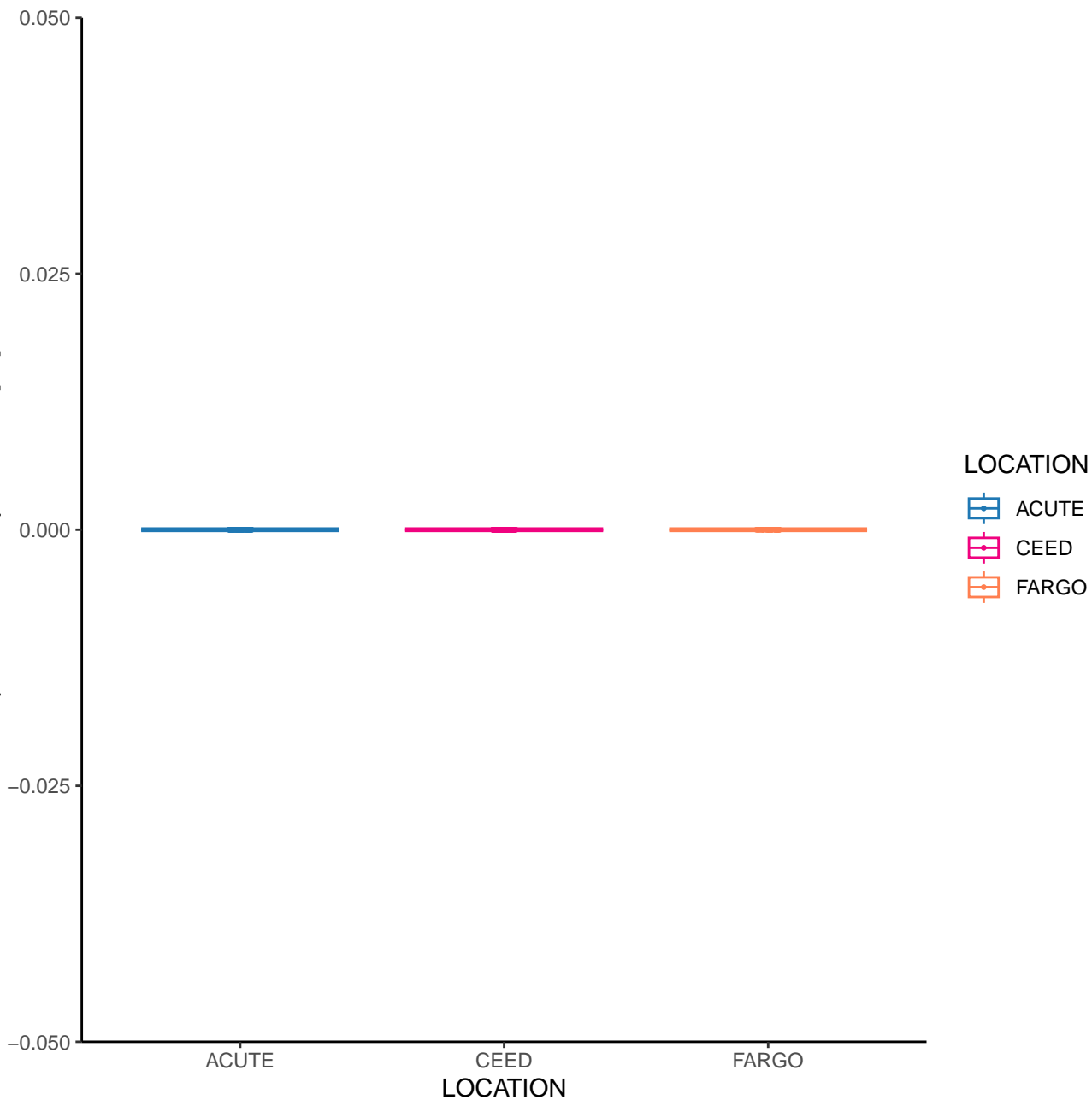
anova p-value: 0.99998

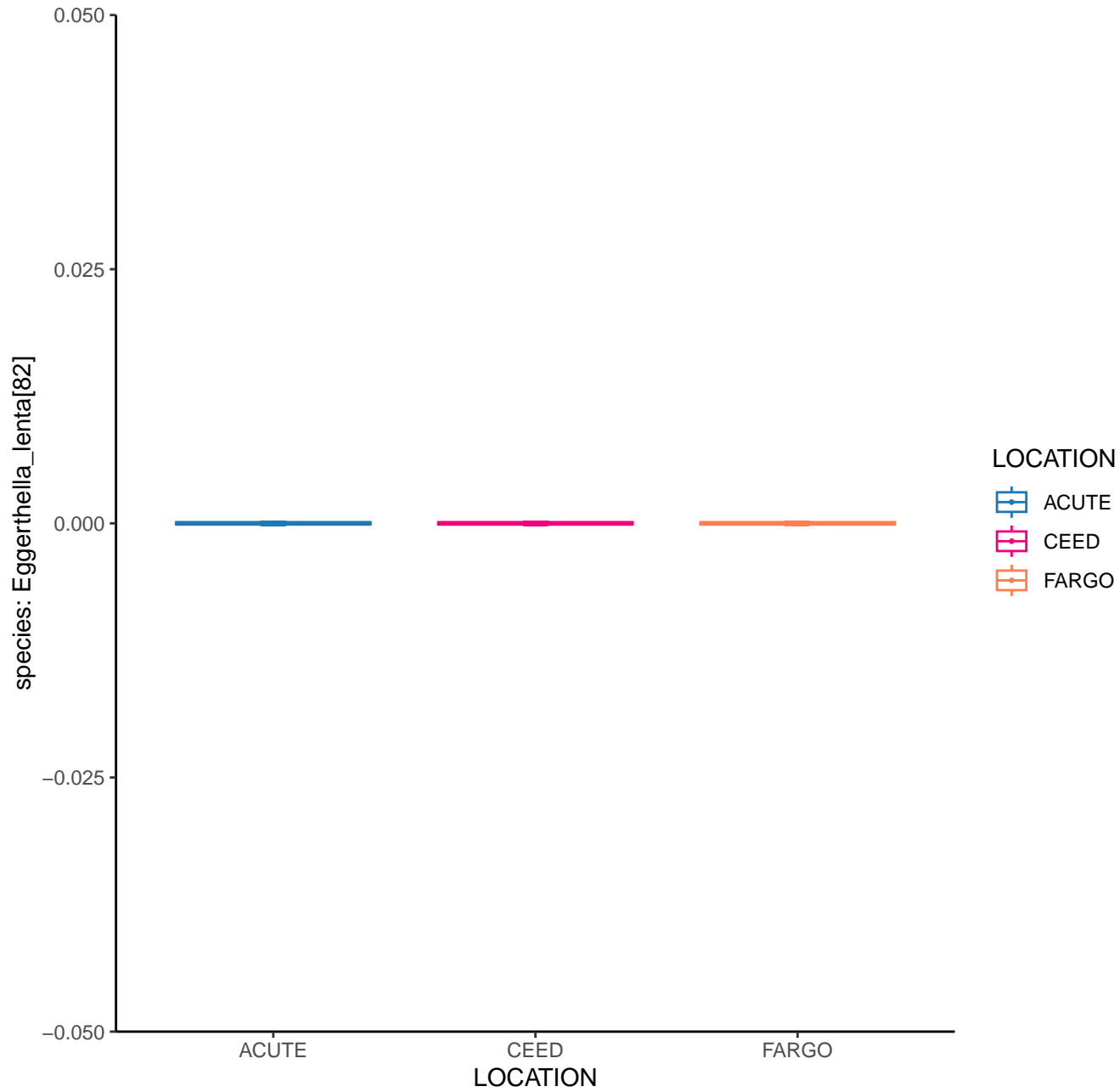
fdr-adjusted p-value: 0.99998

species: g__UCG.003 (UB*)[127]



species: Anaerospirrobacter.[78]





species: *Flavonifractor plautii*[123]

